X65453 M.musculus AF013985 Rattus no AF116582 Rattus no

Sequence Seguence Sequence AR103382 Sequence AR169239 Sequence 187864 Sequence 3 Sequence

AR103367 Sequence AR169224 Sequence IB7865 Sequence 5

AR078308 Sequence

Sequence AR076918 Sequence AR085411 Sequence

AX208160

OM nucleic

Run on:

Sequence:

Searched:

Database

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1 (bases 1 to 818)
Kehry,M. and Castle.B.
Methods for proliferating and differentiating B cells with high
density membrane CD40 ligand
Patent: US 5817516-A 1 06-OCT-1998;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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AR044778
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AR044778
AR171646
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AR085411
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AR171647
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AR044778
                                                                                           29, 2002, 20:42:01; Search time 6499.83 Seconds (without alignments) 2520.909 Million cell updates/sec
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             GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 200000000
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Z15017 H.sapiens m X67878 H.sapiens m

AX090039 Sequence

Sequence 7

Sequence Seguence Human CD40

H.sapiens

AR169232 Sequence AR171647 Sequence

Sequence AR106246 Sequence

AR076926 Sequence Sequence Sequence

AR078316 AR085419 AF344859 Macaca mu X96710 H.sapiens m

248469 B.taurus mR

AF079105 Felis ca

AF086711

AF344860 Aotus tri AF344844 Callithri

PAT 29-SEP-1999

Score 751; DB 6; Length 818;

95.98;

Query Match

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AF344853 Macaca ne AR076929 Sequence AR078319 Sequence

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           Indels
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1 (bases 1 to 918)
Kehry,M. and Castle,B.
Reliculture system comprising high density
  8.9e-185;
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  Pred. No. 8.9e
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Sequence 1 from patent US
AR171646 GI:17910596
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   97.48;
     Similarity
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                                                                                                    Score 751; DB 6;
Pred. No. 8.9e-185;
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Location/Qualifiers
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31-AUG-2000

PAT

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1 (bases 1 to 783)
Armitage, R.Jr., Fanslow, W.C. and Spriggs, M.K.
Monoclonal antibodies to CD40 ligand, pharmaceutical composition
comprising the same and hybridomas producing the same
Patent: US 5961974-A.1 05-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AACAAAGAAGAGAAAAAAGAAAAGCTTTGAAATGCAAAGGGGGGGTGATGAGGATCCTCAA 360
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                              CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT
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Pred. No. 2.3e-184;
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Best Local Similarity 97.3%;
Matches 762; Conservative
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae;
1 (bases 1 to 1250)
Tripp,R.A., Anderson,L.J. and Brown,M.P.
Tato 1 19and adjuvant for respiratory syncytial virus
Patent: WO 0156602-A 1 09-AUG-2001;
The Secretary, Department of Health and Human Service
Location/Qualifiers
                                                                                                                                                                                                                                                                   Length 1250;
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llarity 97.4%; Pred. No. 8.9e-185;
Conservative 0; Mismatches 20;
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Pred. No. 2.3e-184;
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                                               ACGETTAAAAGGAGACTCTATTATGTCTACACTCAAĞTCACCTTCTGCTCTAATCGG
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Unclassified.
Unclassified.
Armitage,R.J., Fanslow,W.C., Spriggs,M.K., Srinivasan,S.
Armitage,R.J., Fanslow,W.C., and McGrew,J.T.
Gibson,M.G., Morris,A.E. and McGrew,J.T.
DNA encoding CD40 ligand, a cytokine that binds CD40
Patent: US 5981724-A.1 pl. NOV-1999;
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Pred. No. 2.3e-184;
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                         CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT
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Armitage,R.J., Fanslow,W.C., Spriggs,M.K.,
Gibson,M.G.
Method of augmenting a vaccine response by
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AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAATGGGAAACAGCTG
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Armitage,R.J., Fanslow,W.C. and Spriggs,M.K.
CD40 ligand polypeptide
Patent: US 6087329-A l 11-JUL-2000;
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Mismatches 21;
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187865.1 GI:3407805
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1 (bases 1 to 783)
Srinivasan, S. and Spri
Methods of preparing s
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              Best Local Similarity 97.3
Matches 762; Conservative
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AUTHORS TITLE REFERENCE

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Daniel K.C., Foss, Y., Moussavi, A., Macary, P., Kemeny, D.M., Farzaneh, F. and Gaken, J.A.
Cloning and sequencing of rat CD40 ligand
Unpublished
2 (bases I to 783)
Daniel, K.C., Foss, Y., Moussavi, A., Macary, P., Kemeny, D.M.,
Farzaneh, F. and Gaken, J.A.
Direct Submission
Submitted (14-JUL-1997) Immunology, King's College School of Medicine and Dentistry, 123 Coldharbourlane, London SES 9NU, United
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ACGGTTAAAAAGAGAAGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGG
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/codon_start=1
/product="CD40 ligand"
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/db_xref="G1:13872517"
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EEKKENSFEMQRGDEDPQIAAHVVSFANSNAASVLQMAKKGYYTMKSNLVMLENGKQL
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                                                                                                                               Armitage, R., Fanslow, W., Sato, T.A., Clifford, K.N., Strockbine, L.,
Macduff, B.M., Anderson, D.M., Gimpel, S.D., Davis-Smith, T.,
Maliszewski, C.R., Clark, B.A., Smith, C.A., Grabstein, K.H., Cosman, D.
                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
1 (bases 1 to 1250)
                                                                                                                                                                                             Molecular and biological characterization of a murine ligand for \mathtt{CD40}
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                                                                                                                                                                                                                                                                                                                                                                                                                                Submitted (24-APR-2001) Strockbine, L. Immunex Research & Development, Molecular Biology, 51 University Street, Seattle, Washington, 98101, USA On Apr 27, 2001 this sequence version replaced gi:50351.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
                                                                                                                                                                                                                                                                                                                 Submitted (07-APR-1992) M.K. Spriggs, Immunex Research & Development, Molecular Biology, 51 University Street, Seattle, Washington, 98101, USA
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Pred. No. 2.3e-184;
0; Mismatches 21;
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/db_xref="taxon:10090"
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                 GI:13872516
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Best Local Similarity 97.3%;
Matches 762; Conservative
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Spriggs, M.K.
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Spriggs, M.K.
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Pred. No. 1.5e-168;
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TVRREGLYYYTOVTFCSNRPPLSORPTIVSIMLKPSSGSERIILIRAANTHSSSKLCE
QOSIHLGGVFELQAGASVVVNTFASQVIHGIGFSSIGLIKL*
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/db_xref="G1:4545250"
/translation="MIETYSOPSPRSVATGLPASMKIFMYLLTTVFLITQMIGSVPFAV
ROD 12-JUL-2000
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, Milwaukee, WI
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Pred. No. 1.1e-167;
0; Mismatches 61;
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     mRNA
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/organism="Rattus norvegicus'
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CD40 ligand mRNA,
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                                                                                                                                                                                                                                                                                                                                        /strain="Sprague Dawley
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/cell_type="splenocyte"
1. .783
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ne rat C
(2000)
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Rodentia;
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Hallett,K.M. and Oaks,M.K.
Nucleotide sequence of the ra
DNA Seq. 10 (6), 405-406 (200
20284949
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Center, 2900 W. Oklahoma Ave.
Location/Qualifiers,
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Hallett, K.M. and Oaks, M.K.
Direct Submission
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                                                GI:4545249
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92.2%;
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Armitage, R.J., Fanslow, W.C. and Spriggs, M.K.
Monoclonal antibodies to C040 ligand, pharmaceutical composition
comprising the same and hybridomas producing the same
Patent: US 5961974-A 22 05-0CT-1999;
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                                                                     GAGCCTTCGAGTCAACGCCCATTCATCGTCGCCTCTGGCTGAAGCCCCAGCATTGGATCT 600
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                                  ACGGTTAAAAGAGAAGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTTCTAATCGG
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Pred. No. 1.2e-153;
0; Mismatches 1;
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Armitage, R.J., Fanslow, W.C., Spriggs, M.K., Srinivasan, S.,
Gibson, M.G., Morris, A.E. and McGrew, J.T.
Recombinant soluble CD40 ligand polypeptide and pharmaceutical
composition containing the same
Patent: US 5962406-A 22 05-OCT-1999;
                                                                                                       644
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TACACTCAAGTCACCTTCTGCTCTAATCGGGAGCCTTCGAGTCAACGCCCATTCATCGTC
                                                                                            GGCCTCTGGCTGAAGCCCAGCATTGGATCTGAGAGAATCTTACTCAAGGCGGCAAATACC
                                                                                                                                                                                CACAGTTCCTCCCCAGCTTTGCGAGCAGCAGTCTGTTCACTTGGGCGGAGTGTTTGAATTA
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Pred. No. 1.2e-153;
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US 5962406.
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                                                                       1 (bases 1 to 878)
Armitege, R.J., Fanslow, W.C., Spriggs, M.K., Srinivasan, S.
Gibson, M.G., Morris, A.E. and McGrew, J.T.
DNA encoding CD40 ligand, a cytokine that binds CD40
Patent: US 5981724-A 22 09-NOV-1999;
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Pred. No. 1.2e-153;
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Sequence 22 from patent US 5981724.
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
                                                                             OM nucleic - nucleic search, using sw model
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Maximum DB seq length: 200000000
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Perfect score:,
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

em_gss_hum:* em_gss_pln:* em_gss_vrt:* em_gss_inv

gb_gss

em_estro:

qb_est1

em_estba:* em_esthum:*

em_estin _estmu _estov em_estpl

ALI03735 Drosophil ALO63921 Drosophil ALO98657 Drosophil AZ691914 ENTMUZGTR ALI00719 Drosophil ALI100719 Drosophil ALS14085 ALS14085 AQ37329 RPCT11-14 ALS14657 ALS14687 B1389827 pppic.pk0 AW486605 75217 MAR AI982044 pat.pk007 AL060767 Drosophil AL534423 AL534423 BF599437 263218 MA AL108993 Drosophil Description CNS0145P CNS0039G CNS0108N AL514657 BI389827 CNS0119V CNS011TX AL514085 AL534423 CNS0187R BF599437 CNS0181N A2691914 AI982044 G Length DB Query Match I Score Š. Result

AZ203738 SP_0084_A AL065906 Drosophil AL237112 Tetraodon AL513719 AL513719 AL100660 Drosophil	AL10881 Drosophil AL514085 AL514085 AQ451757 HS_5184_B AL109264 Drosophil AL188420 Tetraodon A105414 Drosophil	≪.	AD510959 JOSOCHILI AD521025 HS. 5212_B AL513979 AL513979 AG604438 Pan trog1 A155754 Pt. 1-1.A	AL082000 LUSUPULL AL419462 T3 end of AL063926 Drosophil AL098379 Drosophil AL513817 AL513817
AZ203738 CNS006TJ CNS0383B AL513719 CNS011SA	CNS0182P AL514085 AQ451757 CNS018FA CNS02ADN CNS00001	CNS0168L CNS006MT BJ096268 CNS03DHB AL514721 CNS00LLD	CNSUCASW R41111 AQ521025 AL513979 AG044338 AI557564	CNS00055 CNS00039L CNS0100X AL513817
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ALIGNMENTS

	BF599437 492 bp mRNA linear EST 25-APR-2001 26321B MARC 3BOV Bos taurus cDNA 5', mRNA sequence. BF50437			<pre>Smith,T.P.L., Grosse,W.M., Freking,B.A., Roberts,A.J., Stone,R.T., Casas,E., Wray,J.E., White,J., Cho,J., Fahrenkrug,S.C., Bennett ,G.L., Heaton,M.P., Laegreid,W.W., Rohrer,G.A., Chitko-McKown,C.G., Pertea,G., Holt,I., Karamycheva,S., Liang,F., Quackenbush,J. and Keele,J.W.</pre>	Sequence evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle	Genome Res. 11 (4), 626-630 (2001) . 21180013 Contact: Smith TPL	USDA, ARS, US Meat Animal Research Center PO Box 166, Clay Center, NE 68933-0166, USA Tel: 402 762 4366 Fax: 402 762 4396 Fax: 402 762 4396 Email: smith@email.marc.usda.gov	Single pass sequencing. Bases called and alt_trimmed with phred v0.980904.e. Vector identified by cross_match with the -minscore 18 and -minmatch 12 options. PCR PRimers	FORWARD: AGGAAACACTATGACCAT BACKWARD: GTTTTCCCAGTCACGACG Plate: 33 row: N column: 5 Seq primer: ATTTAGGTACACTATAG. Location/Qualifiers e
RESULT 1 BF599437	LOCUS DEFINITION	VERSION KEYWORDS SOURCE ORGANISM	REFERENCE	AUTHORS	TITLE	JOURNAL MEDLINE COMMENT			FEATURES Source

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/db_xref="taxon:9913"
/clone_lib="MARC 1B0V"
/tissue_type="pooled"
/lab_host="DH10B"
                                                                                                                                                         1. .398
/organism="Bos taurus"
                                                                                     FORWARD: AGGAAACAGCTATGACCAT
BACKWARD: GTTTTCCCAGTCACGACG
Plate: 34 row: F column: 23
Seg primer: ATTTAGGTGACACTATAG
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                                                              and -minmatch 12 options.
PCR PRimers
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402 762 4390
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Gallus gallus
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1 (bases 1 to 398)

2 (bases 1 to 398)

Smith, T. P. L., Grosse, M. M., Freking, B. A., Roberts, A. J., Stone, R. T., Grosse, W. W., T. Cho, J., Pahrenkrug, S. C., Bennett, G. L., Heaton, M. P., Laegreid, W. W., Rohrer, G. A., Chitko-McKown, C. G., Pertea, G., Holt, I., Karamycheva, S., Liang, F., Quackenbush, J., and Keele, J. W.
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libraries and construction of a gene index for cattle
Genome Res. 11 (4), 626-630 (2001)
                                       /note-"Vector: pCMV SPORT6; Site_1: XbaI; Site_2: XhoI:
Library made from pooled tissue from marrow, alveolar
macrophage, ovary, fetal semitendonosus muscle, and fetal
longissimus muscle."
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                                                                                                                                                                                                                                                                                                                                                                         TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
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                                                                                                                                                                                                                                                                                                         240
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                                                                                                                                                         DB 10;
                                                                                                                                                         Score 312.2; DB 10;
pred. No. 9.1e-56;
0; Mismatches 73;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                mRNA
                                                                                                        122
                /clone_lib="MARC 3BOV"
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                          /tissue_type="pooled"
/lab_host="DH10B"
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AW486605
                                                                                                           104 g
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82.9%;
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AW486605
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Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
Phasianinae; Gallus.
                                 -minscore 20
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                                                                                                                                                                                                                                                                                                                         /note="Vector: pCMV SPORT6; Site_1: XbaI; Site_2: XhoI; Library made from pooled tissue from lymph node, ovary, fat, hypothalamus, and pituitary."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 241 TIGCIGAACTOTGAGGAGAIGAGAAGGCAATITGAAGACCITGICAAGGAIATAACGITA 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         193 TTACTGAACTGTGAGGAAATTAGAAGCCGGTTTGAAGACTTGGTCAAGGATATAATGCAA 352
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and trimmed with phred
v0.980904.e. Vector identified by cross_match with the -min
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 254.8; DB 9;
pred. No. 1.1e-43;
0; Mismatches 47;
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1 (bases 1 to 797)
Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
Full-length cDNA libraries and normalization
Onpublished (2001)
                                                                                                                                                                            /organism="Drosophila melanogaster"
/db_xref="taxon:7227"
/clone_lib="RPCI-98"
                                                                                                                                                                                                                                                                       538
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Pred. No. 0.07;
l; Mismatches 153;
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/db_xref="taxon:9606"
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1. .797
                                                                                                                                                   Location/Qualifiers
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                                                                                                                                                                                                                                     /clone="BACR12K22"
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AL534423 LTI_FL013_FBrn1
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69; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Web: www.genoscope.cns.fr)
Determination of this BAC-end sequence was carried out as part of collaboration with the Berkeley Drosophila Genome Project (BDGP).
The BDGP is constructing a physicial map of the Drosophila melanogaster genome using these BACs. For further information please see http://www.fruitfly.org The BDGP Drosophila melanogaster BAC library was prepared by Kazutoyo Osoegawa and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Drosophila melanogaster genome survey sequence TET3 end of BAC # BACR12K22 of RPCI-98 library from Drosophila melanogaster (fruit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  671
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    116 GAAGGGACTTGACACGCACACACCACCACCCACTCTGTGACCTCCAGTCCATCCGGGA 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                57 -GCGCCATTCACCCTCTATATTTATTTGTACCTCCCCATGGAAGAGGACCGGCTCCTGAT 115
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      495 AGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGGGAGCCTTCGAGTCA 554
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Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Eukaryota; Neoptera; Endopterygota; Diptera; Brachycera;
"""" Ebhydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              555 ACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCTGAGAGAATCTTACT
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                                                                                                                                                                                                                                                                                                                                                                           Length 638
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /cell_type="Con A-activated splenic T cell"
/lab_host="E.coli TOP10 F/"
/note="Vector: pcDNA3"
171 c 151 g 180 t 4 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  732 CCAAGTGATCCACAGAGTTGGCTTCTCTTTTGGCTTACTCAAACTCT 781
                                                                                                                                                                            /db_xref="taxon:9031"
/clone="pat.pk0072.c9.f"
/clone_lib="chicken activated T cell cDNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
            Molecular Endocrinology
University of Delaware
40 Townsend Hall, Newark, DE 19717, USA
TTE1: 302 831-1345
Fax: 302-831-3411
Email: joan@UDel.Edu, www.chickest.udel.edu
                                                                                                                                                                                                                                                                                                                                                                           Score 72.8; DB 9;
Pred. No. 2e-05;
0; Mismatches 112;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DNA
                                                                                                                                                              /organism="Gallus gallus"
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                                                                                                                                Location/Qualifiers
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Matches 169; Conservative
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797 bp mRNA linear EST 13-FEB-2001 Homo sapiens cDNA clone CSODF004YD24 5 ö oţ Aaron Mammoser in Pieter de Jong's laboratory in the Department of Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPGI-98 and was constructed by partial ECORI digestion of Drosophila DNA provided by the BDGP from the lisogenic strain y2; cn bw sp, the same strain used for the BDGP's Pl and EST libraries. A more detailed description of the library and how to order individual BAC clones, the entire library, or filters for hybridization from the BACPAC Resource Center can be Euteleostomi; ATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCATGAAGATTTTGTATTCATAA 199 320 AAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAAATTGCAGCACACGTTGTAA 379 Gaps 821 RAGAARARARGRRARAARARARRRAGRRRRRARGARGRRRGGRRRGGRRRGRRRARA 260 TGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAGAAGAAAAAAG 641 RRRARGARRRRGRRGRRRRGRARGRRRAGGRRRGRRARGARRGRRRAGRRRRARA Genoscope - Centre National de Sequencage BP 191 91006 EVRY cedex - France Email: seqref@genoscope.cns.fr, Web : www.genoscope.cns.fr. 200 AAAAGCTAAAAGAGATGCAACAAAGGAGAAGGATCTTTATCCTTGCTGAACTGTGAGGAGA GCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCCAAGAAAGGATATTATACCA found at http://bacpac.med.buffalo.edu/drosophila_bac.htm. Primates; Catarrhini; Hominidae; Homo ö 440 TGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTGACGGTTAAAAGA 492 997; Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; others Length Indels

us-08-982-272-3.rst

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Drosophila melanogaster genome survey sequence T7 end of BAC BACNIG11 of DrosBAC library from Drosophila melanogaster (fruit AL), genomic survey sequence.
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Submitted (23-JUL-1999) Genoscope - Centre National de Sequencage :
Submitted (23-JUL-1999) Genoscope - Centre National de Sequencage :
Sp 191 91006 EVRY cedex - FRANCE (E-mail : segref@genoscope.cns.fr
- Web : www.genoscope.cns.fr)
Determination of this BAC-end sequence was carried out as part of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was 'prepared from embryos by Alain Bucheton and Genevieve Payan. It has been constructed in the vector
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                                            when www.genoscope.ons.it.

Determination of this BAC-end sequence was carried out as part of a collaboration with the European Drosophila Genome Project (EGGP) - http://www.edgp.ebi.ac.uk -. This Drosophila melanogaster BAC http://www.edgp.ebi.ac.uk -. This Drosophila melanogaster BAC http://www.edgp.ebi.ac.uk -. This Drosophila melanogaster BAC chirch informed by a lain Billaud at CEPH (Centre d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton and Genevieve Payan. It has been constructed in the vector
23-JUL-1999) Genoscope - Centre National de Sequencage
6 EVRY cedex - FRANCE (E-mail : segref@genoscope.cns.fr
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Genome Project (EDGP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAGAAAACAGCTTTGAAATGCAAAGGGTGATGAGGATCCTCAAATTGCAGCACACGTTG 376
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAGAAGAAAA 316
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           268 WIWIWIWIAAMWCRAAWMGGAWARAAWTRGAWWWRAMMAAAMRAARWAWRAWMAAAWAM 326
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Muscomorpha, Ephydroidea, Drosophilidae, Drosophila.
1 (bases 1 to 1043)
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                                                                                                                                                                                                                                                                                                                                                                       /organism="Drosophila
                                                                                                                                                                                                                                                                                                                                                                                               /plasmid="pBeloBAC11"
/db_xref="taxon:7227"
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                 Submitted (23-JUL-1999)
BP 191-91006 EVRY cedex
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pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 52.8; DB Pred. No. 0.31;
                                /clone="CSODF004YD24"
/clone_lib="LTI_FL013_FBrn1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         36;
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Direct Submission
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Drosophila melanogaster genome survey sequence TET3 end of BAC # BACROSKIO of RPCI-98 library from Drosophila melanogaster (fruit AL063921 GI:4941778
                                                                                                                                                                                                                                                                                                                                                                 ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pl and EST libraries. A more detailed description of the library and how to order individual BAC clones, the entire library, or filters for hybridization from the BACPAC Resource Center can be found at http://bacpac.med.buffalo.edu/drosophila_bac.htm:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      181 GAAGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC 240
                                                                                                                                                                                                                                                                                                                                                                                                                     61 ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           241 TTGCTGAACTGTGAGGAGGATGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          736 ТАТАТАААТТИАААИАТАТАТАТААТАТАТАМАТАААААТАТТТТАТИАИМИМИТААТА 677
                                                                                                                                                                                                                                                                                                                                                                       Gaps
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Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
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                                                                                                                                                                                                                                                                                                          Length 1043;
                                                                                                                                                                                                  167 others
                                                                                                                                                                                                                                                                                                                                                                    Indels
                       /organism="Drosophila melanogaster"
/plasmid="pBeloBAC11"
/db_xref="taxon:7227"
/clone_lib="DrosBAC"
/clone="BACN11G11"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               301 AACAAAGAAGAGAAAAAAGAAAACAGCTTTGAAATGCAAAGA 342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6.3%; SCOIE - 1.4;
35.8%; Pred. No. 1.4;
-*** 48; Mismatches 133;
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                                                                                                                                                                  /note="end
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                                                                                                                                                                                                                                                                                                                                                                 Conservative
.1043
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (23-UUL-1999) Genoscope - Centre National de Sequencage : BP 191 91006 EVRY cedex - FRANCE (E-mail : seqréfégenoscope.cns.fr - Web : www.genoscope.cns.fr)
Determination of this BAC-end sequence was carried out as part of a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LUJB DDNA linear GSS 26-JUL-1: Drosophila melanogaster genome survey sequence SP6 end of BAC BAC MACN03101 of DrosBAC library from Drosophila melanogaster (fruit AL098657
                                                                                                                                                                                                                                                                                                                                                                                                                                                    1084 DDWDWTKWWTWWKDRADDRRWAGDADRWAWDDGAGTWWTATWWWWWWWWATWDTWWDKWW 1025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAAGGAGGAGGATCTTTATCCT 241
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     300 AAACAAAGAAGAGAAAAAAGAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCA 359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGC 419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               420 CAAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCT 479
                                                                                                                                                                                                                                                                                                                                                                                           62 TGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCAC 121
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              724 KRRDTRWDDADADDTARDDRRRRGDDGADAGKGKKTGRKRRRRDRATWDRTDAWWADAAW 665
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1024 WWATAAKTDTAWTWWRTAWRADWAGRDRGAGKRDRDAATDADGAGRRDGGRKRKDKKDKK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DADDDTDGTKDDDDKDKHVDWAKGTWGDATWAWAATDWWWWGWADADWWTWDAAADDW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   122 TITITGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCATG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          964 DGDDDKKGGKKKKAAKAAKWATKWWDDWDKDWKWDGAKDRKADDDDGAGDKDDDGKGK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      242 TGCTGAACTGTGAGGAGTGAGAAGGCA--ATTTGAAGACCTTGTCAAGGATATAACGTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Ins
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
                                                                                                                                                                                                                                                                   Length 1101;
                                                                                                                                        others
                                                                                                                                     503
                                                                                                                                                                                                                                                                   DB 12;
                                                                                                                                     202 t
                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                   Score 48.8;
                                                                                                                                                                                                                                                                                                     Pred. No.
/db_xref="taxon:7227"
/clone_lib="RPCI-98"
/clone="BACR08K10"
                                                                                      /note="end :.TET3"
                                                                                                                                                                                                                                                             6.2%; Sccilarity 13.7%; Pre
Conservative 221;
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Genomic DNA isolated from broth cultures of E. histolytica Genomic DNA isolated from broth cultures of E. histolytica using a method described by Clark and Diamond (Clark, C.G., and Diamond, L.S. (1993) Entamobe histolytica: a c.G., and Diamond, L.S. (1993) Entamobe histolytica: a c.G., and Diamond, L.S. (1993) Entamobe histolytica: a 77:450.) The DNA was mechanically sheared to give a 77:450.) The DNA was mechanically sheared to give a 77:450.) The DNA was mechanically sheared to give a fight size distribution (-2 kb). The v +1 method used for tight size distribution is described in detail in Smith, the library construction is described in detail in Smith, H.O. and Venter, J.C. (Waking small insert libraries for whole genome shotgun sequencing projects. In Genome Sequencing: A Practical Approach, eds. M. Vaudin and B. Barell, Oxford University Press, 1999)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  - Web : www.genoscope.cns.fr)
Determination of this BAC-end sequence was carried out as part of a Determination of this BAC-end sequence was carried out as part of collaboration with the Burogen Drosophila Genome Project (EDGP) - http://www.edgp.ebi.ac.uk - This Drosophila melanogaster BAC library (Dros BAC) was made by Alain Billaud at CEPH (Centre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant: The DNA was prepared from embryos by Alain Bucheton and Genevieve Payan. It has been constructed in the vector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Direct Submission
Submitted (23-JUL-1999) Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - FRANCE (E-mail : segref@genoscope.cns.fr
Clones are derived from the Entamoeba histolytica HM1:IMSS sheared
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         246 GAACTGTGAGGAGATGAGAAGGCAATTTGÅAGACCTTGTCAAGGATATAACGTTAAACÀA 305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Petrygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
I (bases 1 to 1101)
                                                                                                                                                                                                                                                                                                                                                 /clone_lib="Entamoeba histolytica Sheared DNA" /clone_lib="Entamoeba histolytica Sheared DNA" /note="Vector: pHOS1; Site_l: Bst I; Constructed a Institute for Genomic Research (TIGR), Rockville, Rockville, Rockville, Institute for Genomic Research (TIGR), Rockville, Rockvill
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AGAAGAGAAAAAAAAAAAAGCTTTTGAAATGCAAAGAGGTGATGAGGA 353
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ilarity 54.8%; Pred. No. 6.7;
Conservative 0; Mismatches 76;
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                                                                                                                                   High quality sequence start: 15
High quality sequence stop: 330.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         97 ITAATAAAGYRCAANCANCNNTNNNNNNNNNNNCYATAAAATNNTNTTAANAATAAAA 156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTCATAAAAAAGCTAAAGAATGCAACAAAGGAGAAGGATCTTATCCTTGCTGAACTG 251
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   72 TATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCACTTTTTGCTGT 131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            132 GTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCATGAAGATTTTGT 191
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               37 TITINNNITITINNNNNTITICITWANNAACYWITTAITITITITITITATTITITA 96
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                6.1%; Score 47.8; DB 12;
llarity 34.6%; Pred. No. 3.4;
Conservative 20; Mismatches 273;
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Fax: 301 838 9243
Fax: 301 910ftus@tigr.org
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                                                                                                                                             /organism="Drosophila m
/plasmid="pbeloBAC11"
/db_xref="taxon:727"
/clone_lib="DrosBAC"
/clone="BACN03L01"
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Contact: Brendan J Loftus
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Drosophila melanogaster genome survey sequence T7 end of BAC
BACN06A02 of DrosBAC library from Drosophila melanogaster (fruit
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Determination of this BAC-end sequence was carried out as part of a collaboration with the European Drosophila Genome Project (EDGP) http://www.edgp.ebi.ac.uk - This Drosophila melanogaster BAC library (Dros BAC) was made by Alain Billaud at CEPH (Centre
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                                                                             /organism="Drosophila melanogaster"
/plasmid="pBeloBAC11"
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/tissue_type="placenta" // note="Vector: pCMVSPORT 6; Site_1: NotI; lst strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the pCMVSPORT 6
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BP 191 91006 EVRY cedex - France
Email: seqrefégenoscope.cns.fr, Web : www.genoscope.cns.fr.
Location/Qualifiers
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Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
Unpublished (2001)
                                                                                                                                                                                                         others
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                       1. 828
/organism="Drosophila melanogaster"
                                                                                                                                                                                                                                                                                                         ; Score 45.6; DB 12;
; Pred. No. 10;
29; Mismatches 153;
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/db_xref="taxon:7227"
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Location/Qualifiers
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Matches 120; Conservative
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/clone_i.c.
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RPCI11 Human Male AC Library"

165

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BASE COUNT
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(pieter@dejong.med.buffalo.edu). Clones may be purchased from
BACPAC Resources (http://bacpac.med.buffalo.edu/ordering) or from
               Life Technologies. Contact: Feng Liang Life Technologies, a division of Invitrogen 9800 Medical Center Drive Rockville, Maryland 20850, USA Fax: (1) 301 610 8371 Email: fliang@lifetech.com URL: http://fulllength.invitrogen.com.
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http://www.tigr.org/tdb/humgen/bac_end_search/bac_end_search.html
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constructed
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                                                                                                                                                                                                                                                                                                                                                                            128 CTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCATGAAGATT 187
                                                                                                                                                                                                                                                                                                                                                                                                                       244 TITIYITITIAAAGCAAGGCIGGGAAAAAAAANNAAAANNNNAANNNNTAAAAAA 185
                                                                                                                                                                                                                                                                                                                                                                                                                                                             188 ITGTATICATAAAAAGCTAAAGAGATGCAACAAAGGAGGAAGGATCTTTATCCTTGCTGA 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAG 307
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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normalized, Library was
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Shaying Zhao, William Nierman, Mark Adams Department of Bukaryotic Genomics
The Institute for Genomic Research
9712 Medical Center Dr., Rockville, MD 20850
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Zhao, S., Adams, M.D., Nierman, W., Malek, J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       308 AAGAGAAAAAGAAAACAGCTTTGAAATGCAAAGA 342
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          аааааааааааааааааааааааааааааааа 30
                                                                                                                                                                                                                                                          14; Mismatches
                                                                                                                                                                                                                                        Pred. No. 13;
                                                                                                                                                                                                                  Score 45.2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unpublished (1997)
Other_GSSs: RPCI11-14516.TV
      Was
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301 838 0208
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δ qq q οy g δy 8 /organism="Homo sapiens" /db_xref="GDB:7555493"

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was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the pCMVSPORT 6 vector. Library was normalized. Library was constructed by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Life Technologies Contact : Feng Liang Life Technologies, a division of Invitrogen 9800 Medical Center Drive Rockville, Maryland 20850, USA Fax : (1) 301 610 8371 Email : fliang@lifetech.com URL :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               529 bp mRNA linear EST 13-FEB-2001
Homo sapiens CDNA clone CL0BB0022E09 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'PCMVSPORT 6; Site_1: NotI; 1st strand cDNA
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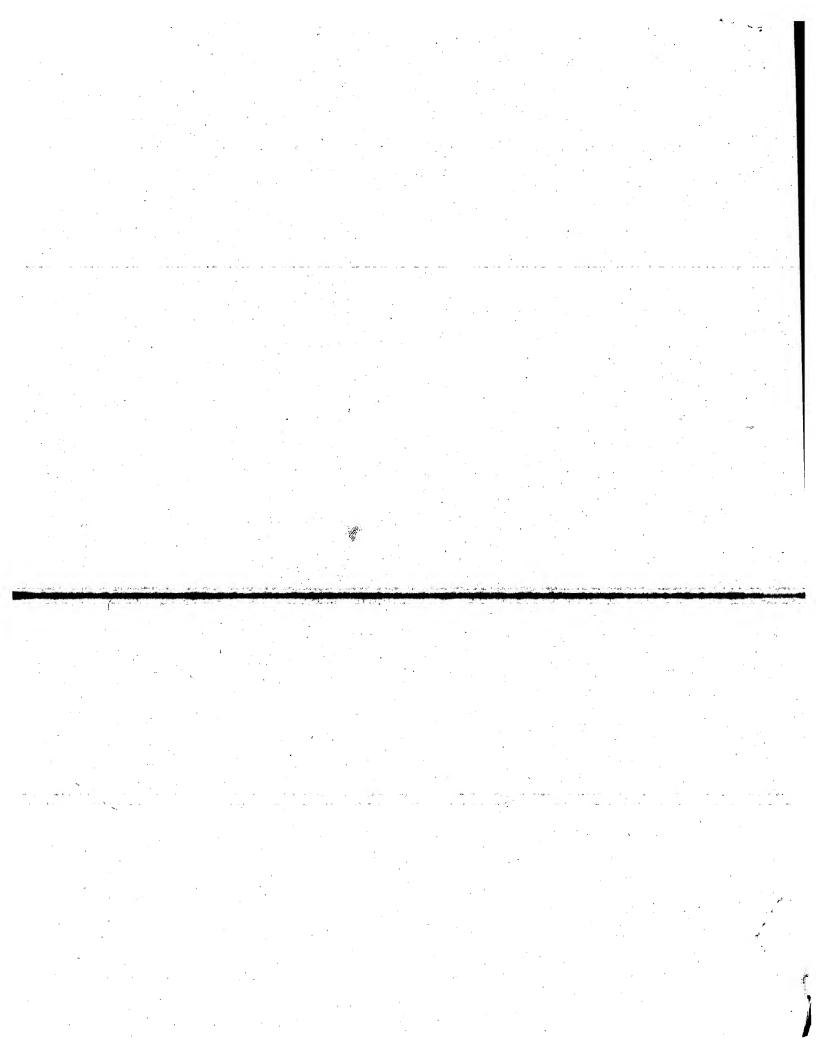
Eukaryota; Metażoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Bammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 529)
                                                                           90 TITICITATCACCCAGATGATTGGGTCAGCACTTTTGCTGTGTATCTTCATAGAAGATT 149
                                                                                                               269 TIGACCIATCICCCAACIGATIGGGITACAGTATATATTIATTATCCACATAAGTACAG 210
                                                                                                                                                     150 GGATAAGGTCGAAGAAGGAAGTAAACCTTCATGAAGATTTTGTATTCATAAAAAAGCTAAA 209
                                                                                                                                                                                           209 TITITAGGIIGAGIGCATAGAAAIGGACAGTGAGCTITAGGAGGCCIATITCACCITIT 150
                                                                                                                                                                                                                                   210 GAGATGCAACAAAGGAGAAGGATCTTTATCCTTGCTGAACTGTGAGGAGATGAGAAGGCA 269
                                                                                                                                                                                                                                                                                                             89 AAAACAAAAACGTGATAAAAGATCAAGGTTAAAATATAGAAGGGATAAAAGGGACAGGAAG 30
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BP 191 91006 EVRY cedex - France
Email: seqref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
                                                                                                                                                                                                                                                                       149 TIGCCCITGAATITCTGGAAGAAGAICTGCTCCTGAAAACAGIATTCTCAAGAGAAGGCC
Length 559;
                                          Indels
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Li,W.B., Gruber,C., Jessee,J. and Polayes,D.
Frul-length cDNA libraries and normalization
Unpublished (2001)
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  DB 12;
                     Pred. No. 13;
0; Mismatches 128;
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/clone_lib="LTI_NFL006_PL2"
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      Score 45.2;
Pred. No. 1
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1. .529
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AL514657
AL514657.1 GI:12778151
        5.8%;
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                                            122; Conservative
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AUTHORS
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AL514657
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	0; Gaps	TG 127	rrr 83	TT 187	TY 143	'GA 247	AA 203	AG 307	 AA 263		
	9; 0;	CTTTI	TTTV	GAAGA	AAAWY	TTGCJ	AMAA	AACA	AAAA?		
47 others	Score 44.6; DB 9; Length 529; Pred. No. 17; ; Mismatches 139; Indels	TTTTTATGRATTTACTTACTGTTTTCTTATCACCCAGATGATTGGGTCAGCACTTTTTG 127	TTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTTWWWTWATTWWTTTWWTTTTTT	128 CTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCATGAAGATT 187	84 TITITWAAATATATATATATAAAWATTTAAAWAAAAAWAATATATAAAAATTTTWAAAAWYYY 143	188 TTGTATTCATAAAAAAGCTAAAGAGGTGCAACAAAGGGGAAGGATCTTTATCCTTGCTGA 247	144 CTTHAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA	248 ACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAG 307	AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA		
	в 9; 139;	ATGAT	WTWAT	GAAGT	TATAT	GAAGG	AAAAA	AAGGA	AAAAA		
89 t	1.6; D 0.17; 1tches	ACCCAG	TTTWW	GAAGAG	AAAWAA	AAAGGA	AAAAAA	сттетс	VAAAAAA	AA 340	W 296
ь Б	5.7%; Score 44.6; 44.3%; Pred. No. 17; tive 13; Mismatches	TTTCTTATC	TTTTTTT	GATAAGGTC	TTTAAAWAA	AGATGCAAC	AAAAAAA	TTTGAAGAC	AAAAAAA	GAAATGCAAA	GCAAAAAAA
20 c	5.7%; 44.3%; ative 13	CTTACTGTT	LTTTTTT	AGAAGATTG	ratatawa Patatawa	AAGCTAAAG	AATTTAAAA	AGAAGGCAA	AAAAAAAA	AACAGCTTT	AAAAARGGG
364 a	Ouery Match 5.7%; Score 44.6; DB 9; Length Best Local Similarity 44.3%; Pred. No. 17; Matches 121; Conservative 13; Mismatches 139; Indels	ATGTATTTAC	TTTTTTT	TATCTTCAT	WAAATATAT	GTATTCATAAAAAAGCTAAAAGAGATGCAACAAAGGAGAAGGATG	AAAAAAA	SAGGAGATG	AAAAAAAA	308 AAGAGAAAAAAGAAAAACAGCTTTGAAATGCAAA 340	264 AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA 296
	tch al Sir 121;	TTTT	TITIL	CTGTG	TTTT	TTGTA	CTTHA	ACTGT	AAAAA	AAGAG	AAAAA
BASE COUNT ORIGIN	Query Match Best Local Matches 12	. 89	24	128	84	188	144	248	204	308	264
BASE CO	Que Bes Mat	γo	ф	οy	QQ	οy	Ω	οy	qq	οy	QQ

Search completed: May 30, 2002, 01:00:01 Job time: 22211 sec



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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd
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OM nucleic - nucleic search, using sw model

May 29, 2002, 23:55:51; Search time 680.87 Seconds Run on:

(without alignments) 1974.450 Million cell updates/sec

Title: Perfect score: Sequence:

US-08-982-272-3 783 1 ATGATCGAAACATACAACCA......TTGGCTTACTCAAACTCTGA 783

IDENTITY_NUC Gapop 10.0 , Gapext 1.0 Scoring table:

1736436 segs, 858457221 residues Searched: Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

/SIDS1/qcqdata/hold-qeneseq/qeneseqn-emb1/NA2002.DAT:

SUMMARIES

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Result No.	Score	Query Match	Duery Match Length DB ID	DB	OI		Description
1 1 1 1			:	-			
-	783	100.0		19	AAV38999		Exemplary CD40 lig
7	773.4	98.8	783	19	AAV39001		Exemplary CD40 liq
æ	751	95.9		19	AAV38998		CD40 ligand gene u
4	751	95.9		19	AAV61062		Murine CD40 ligand
ņ	749.4	95.7	-	15	AAQ63960		Mouse CD40-L type
9	749.4	95.7		19	AAV12853		CD40 ligand coding
7	749.4	95.7		20	AAZ27524		Mouse CD40-L codin
æ	745.2	95.2	782	14	AAQ41507		Murine CD40-L DNA.
σ	648.6	82.8		13	AAV42184	٠	Exemplary nucleoti

Nucleotide sequenc	Mouse trimeric CD4	Human CD40-L type	CD40 ligand gene u				0		Human CD40 ligand		Nucleotide sequenc	Human CD40 ligand	DNA encoding novel	Human CD40L mutein	Genomic sequence o	O-1-(CD40		D40	nd gen	4	CD154	CD154	Feline CD154 cDNA	24	CD154	Canine CD154 cDNA	Canine CD154 cDNA	ectide seg		. HIV-1 gp120 V3 loo	ŗ	. HIV-1 gp120-human	Human CD40-L/Fc fu
AAH25527	AA727538	AA063959	AAV38997	AAV12852	AAQ41506	AA067123	AAT05763	AAT93782	AAV61063	AA227525		AAA51745	AAS86571	AAT58122	AAQ57984	AAQ94091	AAV39000	AAV39002	AAV39003	AAV39004	AAZ55540	AAZ55541	AAZ55542	AAZ55543	AAZ55534	AAZ55535	AA255536	AA255537	AAH25525	AAF82933	AAF82932	AAF82929	AAF82928	AAQ41516
22	200	15	19	19	14	15	16	18	19	20	22	21	23	18	15	16	19	19	19	19	21	21	21	21	21	21	21	77	22	22	22	22	22	14
1477	010	786	786	786	840	840	4	840	840	840	879	1816	81	840	840	840	786	. 98	786	864	882	885	780	780	1878	87	780	78	1552	865	906	2209	25	1425
82.4			75.7	75.7	75.7	75.7	75.7	75.7	75.7	75.7	75.7	75.7				75.0		72.8				68.3						6.99		57.4	57.4	57.4	57.4	57.0
645.2	631 4	592.4	92	592.4	ď	592.4	592.4	592.4	592.4	592.4	592.4	592.4	592.4	591	589.2	587.6	582.8	570	560.4	541.2	535	535	532	532	527	527	524	524	454.6	449.2	•	449.2	449.2	446.2
10	11	13	14	15	16	17	18	19	20	21	22	23	24	25	56	27	28	59	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45
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ALIGNMENTS

AAV38999 standard; DNA; 783 BP AAV38999; AAV38999

RESULT

Exemplary CD40 ligand gene used in the course of the invention:

23-SEP-1998 (first entry)

CD40 ligand; alteration; immunoreactivity; human cell; accessory molecule ligand; AML; gene therapy; treatment; neoplasia; autoimmune disorder; rheumatoid arthritis; vaccine; chimera; ss.

Chimeric - Mus sp. Chimeric - Homo sapiens.

W09826061-A2.

18-JUN-1998

97WO-US22740. 08-DEC-1997;

97US-0982272 96US-0032145 01-DEC-1997; 09-DEC-1996;

(REGC) UNIV CALIFORNIA

Sharma S; Cantwell M, Kipps TJ,

WPI; 1998-348521/30

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The present sequence represents an exemplary CD40 ligand gene, comprising nucleotides encoding the extracellular domains (Domains III and IV) of the murine CD40 ligand gene (AAV38997) operatively linked to uncleotides encoding the extracellular domains (Domains III and IV) and transmembrane domain (Domain II) of human CD40 ligand gene (AAV38998). The sequence is used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of specification describes a method for altering the immunoreactivity of molecule ligand (AML) into the cells so that the AML is expressed on the surface of the calls. Vectors containing the AML is expressed on the surface of the calls. Vectors containing the AML genes can be used in surface of the calls. They can also be used for vaccination to produce freematoid arthritis. They can also be used for vaccination to produce immunity against a virus cell, bacteria, protein, fungus or neoplasia.
vectors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatment of neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
                                                                                                                                        Disclosure; Page 105; 167pp; English
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Sequence 783 BP; 243 A; 161 C; 175 G; 204 T; 0 other;

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                                                                                           CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180
                                                                                                                                            180
                                                                               ATGAAAATTTTTAATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
                                                 Gaps
                                       1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
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  Length 783;
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itch 100.0%; Score 783; DB 19; al Similarity 100.0%; Pred. No: 3.8e-210; 783; Conservative 0; Mismatches 0;
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ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60

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The present sequence represents an exemplary CD40 ligand gene, comprising nucleotides encoding the extracellular domains (Domains III and Iransmembrane domain (Domain II) the murine CD40 ligand gene (AAV38997) operatively linked to nucleotides encoding the cytoplasmic domain of the human CD40 ligand gene (AAV38998). The sequence is used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the introducing a gene encoding an accessory molecule ligand (AML) into the cells of the the AML is expressed on the surface of the cells. Vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 containing the AML genes can be used in gene therapy for treating neoplasia or autoimmune disorders such as rheumatoid arthritis. They also be used for vaccination to produce immunity against a virus cell
                                                  tetgttcaettgggeggagtgtttgaattacaagetggtgettetgtgtttgteaaegtg 720
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                                  ACTGAAGCAAGCCAAGTGATCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC
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Pred, No. 1.9e-207;
); Mismatches 6;
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autoimmune disorder; rheumatoid arthritis; vaccine; ss.
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        gaagattttgtattcataaaaaagctaaagagatgcaacaaaggagaaggatctttatcc
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Pred. No. 3.8e-201;
); Mismatches 20;
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SSXS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence encodes murine CD40 ligand which is used in the method of the invention. The method has been developed for proliferating B cells to increase their number at least 100-fold. The method comprises: (a) providing high density, membrane bound CD40 ligand; and (b) culturing one or more B cells in the presence of this ligand. The culture results in a proliferation in the number of B cells of at least indoced to differentiate into antibody-producing cells in the presence of one or more cytokines. The method as above where the B cells are induced to differentiate into antibody-producing cells in the presence of one or more cytokines. The method can be used for stimulating B-cell of more cytokines and activation is suppressed Eight rounds of the B-cell proliferation and activation is suppressed Eight rounds of division over six days can be achleved, corresponding to a 256-fold
                                                                                                                                       ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCTTTTTGGCTTACTCAAACTC 780
                                                                         TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
 gageettegagteaaegeeeatteategteggeetetggetgaageeeageattggatet 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       by incubating them
                                                                                                                                                                                                                                                                                                                                                                        Murine; mouse; CD40 ligand; TNF receptor family; activated T cell; type 2 membrane glycoprotein; cell proliferation; differentiation; B cell; ds.
                                      GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Increased proliferation of B cells in culture in the presence of membrane-bound CD40 ligand
                                                                                                                                                                                                                                                                                                                                                      Murine CD40 ligand encoding DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
13..795
/*tag= a
/product= "CD40 ligand"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 2; Fig 1; 37pp; English.
                                                                                                                                                                                                                                                                           AAV61062 standard; DNA; 818 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95US-0431055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95US-0431055
94US-0234580
                                                                                                                                                                                                                                                                                                                                 (first entry)
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P-PSDB; AAW71750.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kehry M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           28-APR-1994;
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vast increase compared to previous
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                                                                                                                                          61 ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
                                                                                          1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
                                                                                                       TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTTGTCAACGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAGAGAATCTTACTCAAGGGGGCAAATACCCACAGTTCCTCCCAGCTTGCGAGCAGCAG
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                                                                                                                                                                                                                                                                                                                                                                                                                 GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCT
                                                                                                                                                                                                                                                                                          GAAGATTTTGTATTCATAAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
                                                                                                                                                                                                                       TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA
                                                                                                                                                                                                                                                              ATTGCAGCACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC
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                                                       Length 818;
                                                                          Indels
                             213 T; 0 other;
                                                         ; DB 19;
3.9e-201;
                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              to mRNA; 783 BP.
                                                         Score 751;
Pred. No. 3
                               ö
     18 a
                             C; 186
     numbers, which
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0
                               Sequence 818 BP; 249 A; 170
                                                          Query Match 95.9%;
Best Local Similarity 97.4%;
Matches 763; Conservative
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                culturing methods.
      increase in cell
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540 540 9 900 099 099 720 720

420

us-08-982-272-3.rng

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Leucine zipper; fusion protein production; soluble oligomeric protein; heterologous mammalian type II transmembrane protein; activated T cell; heterologous mammalian type II transmembrane protein; antibody production; CD40-L; B-cell proliferation; CD27-L; lymphocyte antigen; ss.
 aagaaaaggatattataccatgaaaagcaacttggtaatgcttgaaaaatgggaaacagctg 480
                                                                                                                                                                                                                                                                            aacaaagaaagagaaaaagaaaaacagctttgaaatgcaaagaggtgatgatgatcctcaa
                                                                                                                                                       361 attgcagcacacgttgtaagcgaagccaacagtaatgcagcatccgttctacagtgggcc
                                                                                                                                                                                          <u> AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG</u>
                                                                                                                                                                                                                                                                                                                                           GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGGG
                                                                                                                    361 ATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC
                                                                                                                                                                                                                                                          ACGGTTAAAAGAGAAGGACTCTATTATGTCTACACTCCAAGTCACCTTCTGCTCTAATCGG
                                                                                                                                                                                                                                                                                                                         GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATTTTGGCTTACTCAAACTC
                                                                       AAV12853 standard; cDNA to mRNA; 783
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92US-0969703.
93US-0107353.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CD40 ligand coding sequence.
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05-DEC-1991;
23-OCT-1992;
13-AUG-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A DNA fragment encoding the extracellular (soluble) region of mouse CD40-L was ligated to a synthetic oligonucleotide sequence coding for a leader peptide, a 33 amino acid leucine zipper sequence (AAR5968) and the Flag (RTM) linker sequence. Cells expressing the fusion construct are grown to accumulate oligomeric, soluble CD40-L in the supernatant. The leucine zipper sequence spontaneously trimerises in solution and fusion proteins comprising
                                                                                                                                                                                                                    ct= mouse_CD40-L
"nucleotides 148-780 code for the
extracellular region (amino acids 50-260)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 atgatagaaacatacagccaaccttcccccagatccgtggcaactggacttccagcgagc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GAAGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAAGGAGGAGGATCTTTATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  the sequence fused to a heterologous mammalian protein also form
                                                                Leucine zipper; trimerisation; trimeric CD40-L; fusion protein; hetero-oligomer; homo-oligomer; type II transmembrane protein; soluble CD40-L; tumour necrosis factor family; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ó;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 15; Length 783;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ric mammalian proteins - using host protein comprising a leucine zipper
                               Mouse CD40-L type II transmembrane protein coding seguence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 749.4; DB 15;
Pred. No. 1.1e-200;
0; Mismatches 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      cells to express a fusion protein comprising domain and a heterologous mammalian protein
                                                                                                                                                                  Location/Qualifiers
1..783
/*tag a /product= mouse_CD40-/note= "nucleotides 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 1; Page 25-26; 35pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prepn. of soluble oligomeric cells to express a fusion pro
                                                                                                                                                                                                                                                                                                                                                                                      92US-0969703.
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Best Local Similarity 97.3%;
Matches 762; Conservative (
                                                                                                                                                                                                                                                                                                                                                         93WO-US10034
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Srinivasan
                                                                                                                                                                                                                                                                                                                                                                                                                                         (IMMV) IMMUNEX CORP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1994-167465/20.
P-PSDB; AAR53970.
                                                                                                                                    Mus musculus
                                                                                                                                                                                                                                                                                                                                                       20-0CT-1993;
12-JAN-1995
                                                                                                                                                                                                                                                                                                                                                                                                        13-AUG-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Spriggs MK,
                                                                                                                                                                                                                                                                                        WO9410308-A
                                                                                                                                                                                                                                                                                                                                                                                        23-OCT-1992;
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540

(CD40-L)

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The encoded protein can be used in a fusion protein produced using the cathod of the invention. The method is for preparing soluble oligomeric comethod of the invention. The method is for preparing soluble oligomeric protein by culturing a host cell transfected with a vector for the protein by culturing a host cell transfected with a vector for the comparing the cathodran comprises a leucine zipper (used to the W terminus of the extracellular region of a heterologous mammalian type I transmembrane protein or to the C terminus of the mammalian type II transmembrane protein or to the C terminus of the catrodogous mammalian type I transmembrane comparising the leucine zipper is a peptide comprising at least part protein, where the leucine zipper is a peptide comprising at least part of AAW41171 or AAW41172, optionally with conservative amino acid of substitutions, provied that the peptide trimerises in solution. A substitutions, provied that the peptide trimerises in solution. A substitution protein comprising the leucine zipper of AAW41171 linked contributed to a similar manner to membrane-bound CM40-L. A soluble fusion protein a similar manner to membrane-bound CM40-L. A soluble fusion protein a similar manner to membrane-bound CM40-L. A soluble fusion protein comprising the leucine zipper of AAW41172 linked to the extracellular comprising the extracellular region of CD27-L (a type II transmembrane protein the extracellular region of CD27-L (a fusion protein comprising the extracellular region of CD27-L (a fusion protein comprising the extracellular region of CD27-L (a fusion protein comprising the extracellular region of CD27-L (a fusion protein comprising the extracellular region of CD27-L (a fusion protein comprising the extracellular region of CD27-L (a fusion protein comprising the extracellular region of CD27-L (a fusion of CD27-L (a fusion protein comprising the extracellular region of CD27-L (a fusion protein comprising the extracellular region of CD27-L (a fusion protein comprising the extracellular reg
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 241 TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
                                                                                                                                          Soluble oligomeric fusion proteins - comprising leucine zipper fused to extracellular region of transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATÇCGTTCTACAGTGGGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGĠTCGAAGAAGGAAGTAAACCTTCAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GAAGATTTTGTATTCATAAAAAAGCTAAAGAGGTGCAACAAAGGAGGAGGATCTTTATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95.7%; Score 749.4; DB 19; Length 783; nilarity 97.3%; Pred. No. 1.1e-200; Conservative 0; Mismatches 21; Indels 0;
                                                                                                                                                                                                                                                           se is the coding sequence for the mouse CD40 ligand protein can be used in a fusion protein produced us
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 783 BP; 243 A; 159 C; 178 G; 203 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        to EBV-transformed B cells expressing CD27-L.
                                                                                                                                                                                                                             Example 1; column 23-24; 21pp; English
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                                             Spriggs MK, Srinivasan
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Best Local Similarity
Matches 762; Conserv
                                                                                        WPI; 1998-144799/13.
P-PSDB; AAW41179.
(IMMA ) IMMUNEX CORP
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This sequence encodes the mouse CD40 receptor ligand (CD40-L). The invention relates to anti-human CD40-L monoclonal antibodies M90 secreted invention relates to anti-human ED40-L monoclonal antibodies M90 secreted by hybridoma hCD40L-M91 (ATCC HB-12056) and M91 secreted by hybridoma hCD40L-M91 (ATCC HB-12056). M90 and M91 inhibit CD40-L binding to CD40 and the ability of trimeric CD40-L and anti-Immunoglobulin M to induce proliferation of peripheral blood: B cells.
                                                                                                                                         TCTGTTCACTTGGGCGGGGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
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                                                                                                                                                                                                                                                                                                                                                                                                             antibody;
                                                                                                       GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
                                             GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCT
ACGITAAAAGAGAAGICITATTATGICIACACICAAGICACCTICIGGICIAATCGG
                                                                                                                                                                                                                                                                                                                                                                                                             CD40-L; CD40 receptor ligand; anti-human CD40-L monoclonal binding inhibitor; trimeric CD40-L; anti-Immunoglobulin M; peripheral blood B cell; proliferation inhibitor; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anti-human CD40-Ligand monoclonal antibodies
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                                                                                                                                                                                                                                                                                                                        AAZ27524 standard; cDNA; 783 BP.
                                                                                                                                                                                                                                                                                                                                                                                            Mouse CD40-L coding sequence.
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92US-0969703
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23-OCT-1992;
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Length 7,83

DB 20;

Score 749.4;

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Query Match

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                            CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT
                                                                                                                       gaagattttgtattcataaaaagctaaagagatgcaacaaaggagaaggatctttatcc
                                                                                                                                                                                 aacaaagaagagaaaaaagaaaacagctttgaaatgcaaagaggtgatgaggatcctcaa
                                                                                                                                                                                                                          AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG
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                                                                                                                                    TTGCTGAACTGTGAGGAGGTGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA
                                                                                                                                                                  AACAAAGAAGAAAAAAAAAAAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAA
                                                                                                                                                                                             ATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC
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                                                                                                                                                                                                                                                                                                            GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
                                                                                                                                                                                                                                                                                                                     transmembrane; region; intracellular; soluble; activity; B cell; proliferation; induction; antibody; secretion; IgE; agonist;
        Indels
Pred. No. 1.1e-200;
; Mismatches 21;
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97.3%;
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This sequence encodes a murine CD40-L polypeptide which binds to CD40. CD40-L is a type II membrane polypeptide which has an extracellular region at its C-terminus, a transmembrane region and an intracellular region at its N-terminus. A soluble form of CD40-L lacks the transmembrane domain. CD40-L activity is mediated by binding with CD40 and induces B cell proliferation and induction of antibody secretion, including IgE. Membrane bound CD40-L acts as a CD40 agonist and soluble CD40-L acts as a CD40 and and an abinding assay to detect cells expressing CD40-L
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graft-versus-host disease and insulin-dependent diabetes mellitus
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Pred. No. 1.6e-199;
); Mismatches 23;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        cytokine CD40-L as CD40 agonist and
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antagonist; binding assay;
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Matches 759; Conservative
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SSSSS
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aagaaaggatattataccatgaaaagcaacttggtaatgcttgaaaatggggaaacagctg 480
                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; alteration; immunoreactivity; human cell; accessory molecule ligand; AML; gene therapy; treatment; neoplasia; autoimmune disorder; rheumatoid arthritis; vaccine; chimera; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                of
                       GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
                                                                                                                               ACTGAAGCAAGCCAAGTGCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC
                                                                      GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCCAGCATTGGATCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    es - used for
for treatment
                                                                                                                                                                                                                                                                                                                                                                                                                Exemplary nucleotide sequence of a chimeric accessory molecule.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Vectors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatmen neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
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96US-0032145
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Chimeric - Mus sp.
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autoimmune disorders such as rheumatoid arthritis. They can also used for vaccination to produce immunity against a virus cell, bacteria, protein, fungus or neoplasia.
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                                                                Length 783;
                                                                                      Indels
                                     Sequence 783 BP; 252 A; 167 C; 166 G; 198 T; 0 other;
                                                                  DB 19;
                                                                                      84;
                                                                              .4e-172;
                                                                   Score 648.6; Dered. No. 2.4e-0; Mismatches.
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AGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAGAAGAGA 313

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murine surfactant protein of including hub region, collagenous portion, and neck) fused to the murine CD40 ligand extracellular region (including stalk). The specification describes a method for constructing stable bloactive fusion proteins of the difficult to express tumor necrosis factor superfamily (TNFSF) proteins (especially CD40 ligand) as multimeric ligands fused onto branched protein backbones such as collectin molecules e.g. pulmonary surfactant protein D (SPD). The fusion proteins of the invention are useful for stimulating immune response in potentially immunocompetant cells (e.g., resting B cells). They are also useful for increasing antigenicity of cells such as tumor cells or human immunodeficiency virus (HIV) positive cells. They are also useful as a vaccine adjuvant since they stimulate B cells, makes them else likely to diffuse into the circulation, they can be advantageously
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               used as a vaccine adjuvant or tumor immunotherapy agent, injected locally to prevent them from diffusing away. Also, the TNFSF-collectin fusion proteins present new possibilities for the expression of highly active,
                                                                                                                                                                                                                                                                                                                           /note- "signal peptide from murine surfactant protein D"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            multimeric, soluble TNFSF members. CD40L was a powerful stimulant for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Producing tumor necrosis factor superfamily proteins as multimeric ligands fused onto collectin molecules e.g. pulmonary surfactant protein D, useful as vaccine adjuvants against infectious agents and
                                                                                             Fusion protein; CD40 ligand; tumour necrosis factor; TNF; collectin; pulmonary surfactant protein D; SPD; immunocompetant cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      the present sequence encodes a fusion protein comprising the mature
                                                         Nucleotide sequence of surfactant protein D fused to CD40 ligand.
                                                                                                                  immunocompetant cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1477 BP; 418 A; 339 C; 421 G; 299 T; 0 other;
                                                                                                                  pulmonary surfactant protein D; sru; immunocompercell antigenicity; vaccine adjuvant; chimera; ss.
                                                                                                                                                                                                                                  Location/Qualifiers 32..1444 /*tag= a 32..88
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Chimeric - Mus sp.
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aaaaagaaaaacagctttgaaatgcaaagaggtgatgagggtcctcaaattgcagcacacg 1034
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                                                                                                                   ATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTGACGGTTAAAAGAG 493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      High density membrane-bound CD40 ligand; B-lymphocyte; B-cell; differentiation; proliferation; baculovirus; Spodoptera frugiperda; Sf9; insect cell culture; tumour necrosis factor receptor; ss.
AAAAAGAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAAATTGCAGCACACG
                                                         TTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCCCAAGAAAGGATATT
                                                                                                                                                                              AAGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGGGAGCCTTCGAGTC
                                                                                                                                                                                                                                           AACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCTGAGAGAATCTTAC
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13..795
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Murine CD40 ligand DNA.
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Indels

Length 1477;

DB 22;

82.4%; Score 645.2; DB 22; 99.5%; Pred. No. 2.8e-171; ive 0; Mismatches 3;

Conservative

647;

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Local Similarity

Query Match Matches 134 ATCTTCATAGAAGATTGGATAAGGTCGAAGAAGAAGTAAACCTTCATGAAGATTTTGTAT 193

795 atggccatagaagattggataaggtcgaagaggaagtaaaccttcatgaagattttgtat 854

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ATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTGTACAGTGGGCC 420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG 480
                                                                                                encodes a mouse high-density,
                                                                                                                                                                                                                                                                                                                                                                                                          GAAGATTTTGTATTCATAAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 aacaaagaagagaaaaaaaaaaaagaagctttgaaatgcaaagaggtgatgaggatcctcaa
                                       producing
                                                                                   The nucleotide sequence given in AAT05762 encodes a mouse high-dens membrane-bound (hdmb) CD40 ligand (AAR85485) that induces long-term proliferation of B-cells in culture. These proliferating B-cells can be induced to differentiate into antibody-prodg cells. The nucleotide sequence is incorporated into a baculovirus vector that is used to transfect sf9 insect cells for prodn. of recombinant hdmbCD40.
                                                                                                                                                                                                                                                                      ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
                                                                                                                                                                                                                                                                                                                                                             CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAAGAAGTAAACCTTCAT
                                                                                                                                                                                                                                                  ;
0
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                            High density membrane bound CD40 ligand - for stimulating proliferation of B cells in vitro or in vivo, partic. for differentiated cells
                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                           Sequence 818 BP; 237 A; 182 C; 188 G; 211 T; 0 other;
                                                                                                                                                                                                                              Score 632.6; DB 16;
Pred. No..7.7e-168;
0; Mismatches 94;
                                                                                                                                                                                                                                                      Mismatches
                                                                             Disclosure, Fig 1; 74pp; English.
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llarity 88.0%;
Conservative
WPI; 1995-393038/50.
P-PSDB; AAR85485.
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689; Conserv
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This sequence encodes the mouse trimeric CD40 receptor ligand (CD40-L). The invention relates to anti-human (CD40-L monoclonal antibodies M90 secreted by hybridoma hCD40L-M90 (ATCC HB-12055) and M91 secreted by hybridoma hCD40L-M90 (ATCC HB-12055). M90 and M91 inhibit CD40-L binding hybridoma hCD40L-M91 (ATCC HB-12056). M90 and M91 inhibit CD40-L binding to CD40 and the abllity of trimeric CD40-L and anti-Immunoglobulin M to induce proliferation of peripheral blood B cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 878;
                                                                                                                                                                                                                CD40-L; CD40 receptor ligand; anti-human CD40-L monoclonal binding inhibitor; trimeric CD40-L; anti-Immunoglobulin M; peripheral blood B cell; proliferation inhibitor; ss.
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Pred. No. 1.7e-167;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Anti-human CD40-Ligand monoclonal antibodies
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                                                                                                                                                                                             Mouse trimeric CD40-L coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                Armitage
                                                                                                                         standard; cDNA; .878 BP
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illarity 99.8%;
Conservative 0
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91US-0805723.
92US-0969703.
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                                                                                                                                                                           (first entry)
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nes 632; Conserv
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23-OCT-1992;
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"nucleotides 148-783 code for the
extracellular region (amino acids 50-261)"
                                                                                                                                                                                                                                                                                                                                                                                                                TACACTCAAGTCACCTTCTGCTCTAATCGGGAGCCTTCGAGTCAACGCCCATTCATCGTC
                                                                                                                                                                                            GCCCTCTGGCTGAAGCCCAGCATTGGATCTGAGAGAATCTTACTCAAGGCGGCAAATACC
                                                     TTGGTAATGCTTGAAAATGGGAAACAGCTGACGGTTAAAAGAGAAGGACTCTATTATGTC
                                   AGTAATGCAGCATCCGTTCTACAGTGGGCCAAGAAAGGATATTATACCATGAAAAGCAAC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Leucine zipper; trimerisation; trimeric CD40-L; fusion protein; hetero-oligomer; homo-oligomer; type II transmembrane protein; soluble CD40-L; tumour necrosis factor family; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prepn. of soluble oligomeric mammalian proteins - using host cells to express a fusion protein comprising a leucine zipper domain and a heterologous mammalian protein
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P-PSDB; AAR53969.
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13-AUG-1993;
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                                   A DNA fragment encoding the extracellular (soluble) region of human CD40-L was ligated to a synthetic oligonuclectide sequence coding for a leader peptide, a 33 amino acid leucine zipper sequence (AAR53968) and the Flag (RTM) linker sequence. Cells expressing the fusion construct are grown to accumulate oligomeric, soluble CD40-L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA
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                                                                                                                                                                                 the sequence fused to a heterologous mammalian protein, also form
                                                                                                                                                                                                                                                                                                                                                  3;
                                                                                                                                                                                                                                                                                                      Length 786;
                                                                                                                                         supernatant. The leucine zipper sequence spontaneously ses in solution and fusion proteins comprising
                                                                                                                                                                                                                                                                                                                                                  Indels
                                                                                                                                                                                                                                               Sequence 786 BP; 250 A; 168 C; 168 G; 200 T; 0 other;
                                                                                                                                                                                                                                                                                                          DB 15;
                                                                                                                                                                                                                                                                                                        Score 592.4; DB 15;
Pred. No. 1.5e-156;
); Mismatches 111;
  English
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22-23; 35pp;
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illarity 85.5%;
Conservative
                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
  Page
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Leucine zipper; fusion protein production; soluble oligomeric protein; heterologous mammalian type II transmembrane protein; activated T cell; heterologous mammalian type I transmembrane protein; antibody production; CD40-L; B-cell proliferation; CD27-L; lymphocyte antigen; ss.
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                                                                             CTGACGGTTAAAAGAGAAGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAAT
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23-OCT-1992;
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                                                                                                                                                                                                                CD40 ligand; alteration; immunoreactivity; human cell; accessory molecule ligand; AML; gene therapy; treatment; autoimmune disorder; rheumatoid arthritis; vaccine; ss.
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                                                                                                                                                                                    CD40 ligand gene used in the course of the invention.
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luble oligomeric fusion proteins – comprising leucine zipper fused extracellular region of transmembrane protein Example 1; column 19-20; 21pp; English 93US-0107353 ŝ Srinivasan (IMMV) IMMUNEX CORP. WPI; 1998-144799/13. P-PSDB; AAW41178. 13-AUG-1993; Spriggs MK, Soluble 9

The encoded protein can be used in a fusion protein produced using the method is for preparing soluble oligomeric protein by culturing a host call transfected with a vector for the fusion protein. The soluble oligomeric proteins comprise a leucine zipper fused to the N terminus of the extracellular region of a heterologous mammalian type II transmembrane protein or to the C terminus of the extracellular region of a heterologous mammalian type I transmembrane protein or to the C terminus of the extracellular region of a heterologous mammalian type I transmembrane protein, where the leucine zipper is a peptide comprising at least part of AAW4111 or AAW41172, optionally with conservative amino acid substitutions, provided that the peptide trimerises in solution. A soluble fusion protein comprising the leucine zipper of AAW41171 linked to the extracellular region of CD40-L (a type II transmembrane protein antiper is found on activated T cells and acts as a ligand for the B-cell antiper CD40) stimulates B-cell proliferation and antibody production in a similar manner to membrane-bound CD40-L. A soluble fusion protein sequence is the coding sequence for the human CD40 ligand (CD40-L) antigen CD27) inhibits binding of CD27c (a fusion protein the extracellular region of CD27 and a human IgG1 Fc region) comprising the leucine zipper of AAW41172 linked to the extracellular region of CD27-L (a type II transmembrane protein that binds to the lymphocyte antiqen CD27) inhibits binding of CD27c (a fusion protein EBV-transformed B cells expressing CD27-L comprising

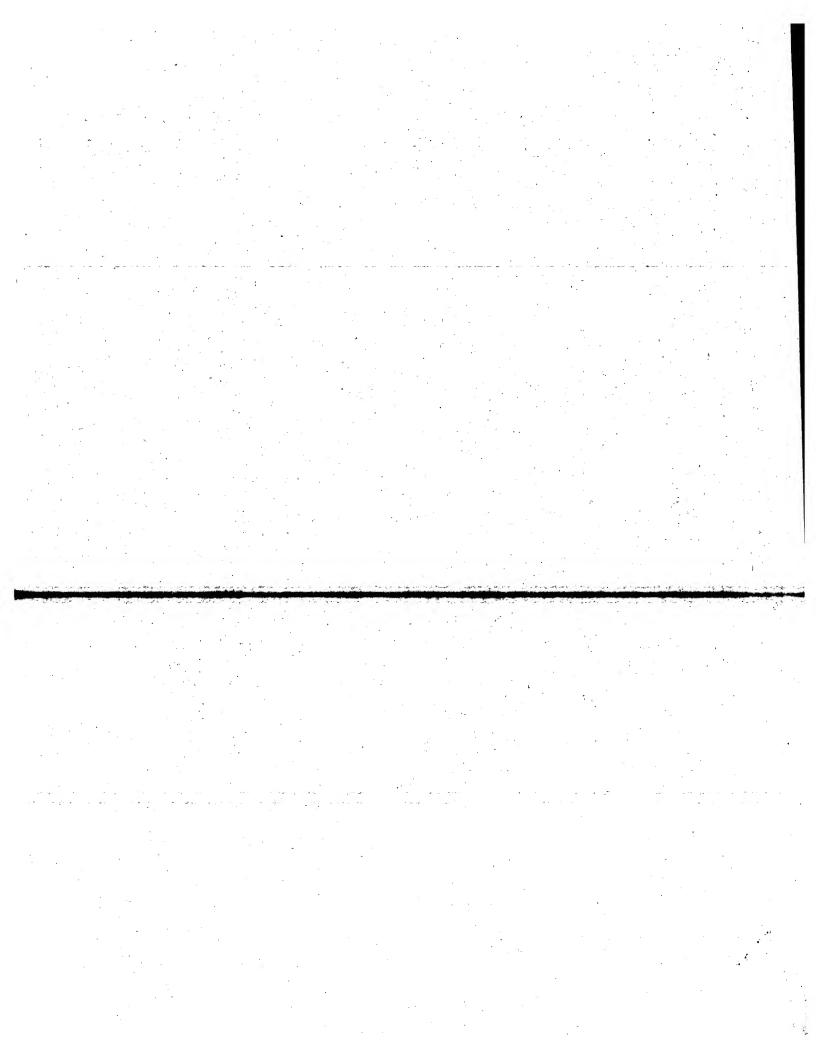
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300 CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180 GAAGATTTTGTATTCATAAAAAGCTAAAGGGAGATGCAACAAAGGAGAAGGATCTTTATCC 240 CAAATTGCAGCACATGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGG 417 ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120 AACAAAGAAGA - - - GAAAAAAAAAAAAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCT 357 1 ATGATCGAAACATACAACCATCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60 TTCCTGAACTGTGAGGAGGTGAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA DB 19; Length 786; Indels Score 592.4; DB 19; Pred. No. 1.5e-156; 0; Mismatches 111; Query Match 75.7%; Best Local Similarity 85.5%; Matches 672; Conservative 301 19 121 241 358 361 418 121 181 181 241 301 61 g g ò g g δ ద ò ò ô ò ò 셤 ò

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SUMMARIES

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	4 80.6 878 17 US-09-392-618-22 Sequence 2 4 80.6 878 17 US-09:399-106-22 Sequence 2 4 75.7 786 5 US-08-107-353-33 Sequence 3, 4 75.7 786 13 US-08-982-272-1 Sequence 1, 4 75.7 786 13 US-08-982-272-1 Sequence 0.1 1.5.7	75.7 840 1 PCT-US94-10780 5 Sequence 75.7 840 1 PCT-US97-11956-1 Sequence 75.7 840 3 US-07-805-723-10 Sequence 75.7 840 3 US-07-805-723-10 Sequence 75.7 840 3 US-07-966-7031-11 Sequence 75.7 840 3 US-07-966-7031-11 Sequence	75.7 840 3 US-07-969-703B-11 Sequence 75.7 840 4 US-08-009-258-7 Sequence 75.7 840 5 US-08-172-664-1 Sequence 75.7 840 6 US-08-172-664-1 Sequence	75.7 840 6 US-08-234-580-3 Sequence		72-3 72-3 INPORMATION: ANT: Kipps, Thomas J.	Cantwell, VENTION: VENTION:	DE INVENTION: DE INVENTION: DE INVENTION:	OF S ONDE	Q U >	: 90/1/1-20RM: IUM TYPE: 3.5" Diskette, UUM TYPE: 144 Mb Storage	0.0	LICATION NUMBER: US/US/US/US/US/US/US/US/US/US/US/US/US/U	ING DATE: 12/9/96 ING DATE: 12/9/96 INFYAGENT INFORMATION: E: Guise, Jeffrey W.	-	EPHONE: (213) 489-1600 EFAX: (213) 955-0440 EX: 67-3510	w	E: nucleic acid ANDEDNESS: single 02.0047: linear	100.0%; Score 783; DB 13; Length 7 Similarity 100.0%; Pred. No. 3.6e-186;
	4 80.6 878 17 US-09-392-618-22 Sequence 2 4 80.6 878 17 US-09:399-106-22 Sequence 2 4 75.7 786 5 US-08-107-353-3 Sequence 3, 4 75.7 786 13 US-08-982-272-1 Sequence 1, 4 75.7 786 13 US-08-982-272-1 Sequence 0.1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	75.7 840 1 PCT-US94-10780 5 Sequence 75.7 840 1 PCT-US97-11956-1 Sequence 75.7 840 3 US-07-805-723-10 Sequence 75.7 840 3 US-07-805-723-10 Sequence 75.7 840 3 US-07-966-7031-11 Sequence 75.7 840 3 US-07-966-7031-11 Sequence	840 3 US-07-969-703B-11 Sequence 840 4 US-08-009-258-7 Sequence 840 5 US-08-172-664-1 Sequence 840 f IIS-0R-215-862-1 Sequence	75.7 840 6 US-08-234-580-3 Sequence		2-2723 ce 3, Application US/08982272 AL INFORMATION: LICANT: Kipps, Thomas J.	Cantwell, VENTION: VENTION:	OF INVENTION: OF INVENTION: OF INVENTION:	OF S ONDE	Q U >	ALTH: 90/1.ZOV. AVOIDED FORM: REMANBLE FORM: REDIUM TYPE: 3.5" Diskette, REDIUM TYPE: 1.4 Mb storage	0.0	APPLICATION NUMBER: US/U8/392.2/2 CLASSIFICATION: LOR APPLICATION ADD TANTON NUMBER: 06/132145	AFFLICITOR DATE: 12/9/96 TORNEY/AGENT INFORMATION: NAME: Guise, Jeffrey W.	-	TELEPHONE: (213) 489-1600 TELEFAX: (213) 955-0440 TELEX: 67-3510	w	TYPE: nucleic acid STRANDINESS: single TOPOLOGY: linear 082-272-3	100.0%; Score 783; DB 13; Length 7 Similarity 100.0%; Pred. No. 3.6e-186;
	631.4 80.6 878 17 US-09-392-618-22 Sequence 2 631.4 80.6 878 17 US-09:399-106-22 Sequence 2 592.4 75.7 786 5 US-08-107-353-3 Sequence 3, 252.4 75.7 786 13 US-08-982-272-1 Sequence 1.	592.4 75.7 840 1 PCT-US94-00780 7 Sequence 592.4 75.7 840 1 PCT-US97-11956-1 Sequence 592.4 75.7 840 1 PCT-US97-11956-1 Sequence 592.4 75.7 840 3 US-07-805-723-10 Sequence 503.4 75.7 840 3 US-07-805-723-10 Sequence	592.4 75.7 840 3 US-07-969-703B-11 Sequence 592.4 75.7 840 4 US-08-099-258-7 Sequence 592.4 75.7 840 5 US-08-08-12-664-1 Sequence 592.4 75.7 840 6 US-08-15-862-1 Sequence	75.7 840 6 US-08-234-580-3 Sequence	1	-982-272-3, Application US/08982272 uence 3, Application US/08982272 NERAL INFORMATION: APPLICANT: Kipps, Thomas J.	Cantwell, VENTION: VENTION:	DE INVENTION: DE INVENTION: DE INVENTION:	SEE	E: LO	FORM: 5% Diskette, 44 Mb storag	0.0	APPLICATION NUMBER: US/US/392/2/2 FILING DATE: CLASSIFICATION: ADDITCATION DATA: ADDITCATION NUMBER: 60/132145	35.50	34,613 SER: 231 RMATION:	TELEPHONE: (213) 489-1600 TELEFAX: (213) 955-0440 TELEX: 67-3510	w	TYPE: nu STRANDEDN TOPOLOGY:	100.0%; Score 783; DB 13; Length 7 Similarity 100.0%; Pred. No. 3.6e-186;
	631.4 80.6 878 17 US-09-392-618-22 Sequence 2 631.4 80.6 878 17 US-09:399-106-22 Sequence 2 592.4 75.7 786 5 US-08-107-353-3 Sequence 3, 252.4 75.7 786 13 US-08-982-272-1 Sequence 1.	75.7 840 1 PCT-US94-10780 5 Sequence 75.7 840 1 PCT-US97-11956-1 Sequence 75.7 840 3 US-07-805-723-10 Sequence 75.7 840 3 US-07-805-723-10 Sequence 75.7 840 3 US-07-966-7031-11 Sequence 75.7 840 3 US-07-966-7031-11 Sequence	592.4 75.7 840 3 US-07-969-703B-11 Sequence 592.4 75.7 840 4 US-08-009-258-7 Sequence 592.4 75.7 840 5 US-08-08-12-664-1 Sequence 592.4 75.7 840 6 US-08-15-862-1 Sequence	75.7 840 6 US-08-234-580-3 Sequence	1	-08-982-272-3 Sequence 3, Application US/08982272 GENERAL INFORMATION: APPLICANT: Kipps, Thomas J.	CANT: SHAIMA, STANT: CANTWELL, OF INVENTION:	OF INVENTION: OF INVENTION: OF INVENTION:	OF S ONDE	Q U >	COMPUTER READABLE FORM: WEDIUM TYPE: 3.5" Diskette, MEDIUM TYPE: 1.44 Mb storage	0.0	APPLICATION NUMBER: US/US/961/1/2 FILING DATE: US/US/961/1/2 FILING DATE: CLASSIFICATION: PRIOR APPLICATION DATA: ADDITORATION NUMBER: 60/132145	FILING DATE: 12/9/96 ATTORNEY AGENT INFORMATION: NAME: Guise, Jeffrey W.	-	TELEPHONE: (213) 489-1600 TELEFAX: (213) 955-0440 TELEX: 67-3510	CHARACTERISTICS: 783 base pairs	TYPE: nu STRANDEDN TOPOLOGY:	Score 783; DB 13; Length 7 Pred. No. 3.6e-186;
	631.4 80.6 878 17 US-09-392-618-22 Sequence 2 631.4 80.6 878 17 US-09:399-106-22 Sequence 2 592.4 75.7 786 5 US-08-107-353-3 Sequence 3, 252.4 75.7 786 13 US-08-982-272-1 Sequence 1.	592.4 75.7 840 1 PCT-US94-00780 7 Sequence 592.4 75.7 840 1 PCT-US97-11956-1 Sequence 592.4 75.7 840 1 PCT-US97-11956-1 Sequence 592.4 75.7 840 3 US-07-805-723-10 Sequence 503.4 75.7 840 3 US-07-805-723-10 Sequence	592.4 75.7 840 3 US-07-969-703B-11 Sequence 592.4 75.7 840 4 US-08-009-258-7 Sequence 592.4 75.7 840 5 US-08-08-12-664-1 Sequence 592.4 75.7 840 6 US-08-15-862-1 Sequence	75.7 840 6 US-08-234-580-3 Sequence		982-272-3 ence 3, Application US/08 ERAL INFORMATION: PPLICANT: Kipps, Thomas	Cantwell, VENTION: VENTION:	OF INVENTION: OF INVENTION: OF INVENTION:	OF S ONDE	Q U >	COMPUTER READABLE FORM: MEDIUM TYPE: 3.5" Diskette, MEDIUM TYPE: 144 Mb storage	0.0	PILING DATE: CLASSIFICATION PRIOR APPLICATION DATA: ADDITICATION ADDIT	FILING DATE: 12/9/96 ATTORNEY/AGENT INFORMATION: NAME: Guise, Jeffrey W.	-	; TELEPHONE: (213) 489-1600 ; TELEFAX: (213) 955-0440 ; TELEX: 67-3510	w	TYPE: nucleic acid STRANDEDNESS: single TOPOLGY: linear US-08-982-272-3	100.0%; Score 783; DB 13; Length 7 Similarity 100.0%; Pred. No. 3.6e-186;

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661 TCTGTTCACȚTGGGCGGAGTGTTTGAĄTTACAAGCTGGTGCTTCTGTGTTTTGTCAACGTG 720
                                                                                                                                                                                                                                                                                         GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGGTTTGCGAGCAGCAGCAG
           541 GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCT
                                                                                                                                                                                                    TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTION Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
RPLICATION NUMBER: US/07/969,703B
FILING DATE: 19921023
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: ARMITACE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
TITLE OF INVENTION: NOYEL CYTOKINE
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 3
US-07-969-703B-1
; Sequence 1, Application US/07969703B
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 206587Ubuo
INPORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 199210:
CLASSIFICATION: '435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: SEATTLE
STATE: WASHINGTON
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CLONE: CD40-L
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                                                                                                                                                               OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: FESTSEQ Version 2.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/982,272 FILING DATE:
                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNBER: 60/132145
FILING DATE: 12/9/96
ATTORNEY/AGENT INFORMATION:
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                IBM Compatible
                                                                                                                                                                                                                                                                                                                                                         NAME: Guise, Jeffrey W. REGISTRATION NUMBER: 34,613
                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: 1.44 Mb storag
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TELEPHONE: (213) 489-1600
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Best Local Similarity 99.2%;
Matches 777; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                      Los Angeles
California
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                                                                          COUNTRY: U.S.A. ZIP: 90071-2066
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CURRENT APPLICATION DATA:
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                                       NUMBER OF SEQUENCES:
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California
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Best Local Similarity
Matches 763; Conserv
                                                                                                         RY: U.S.A.
90071-2066
                              INVENTION:
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                               95.9%; Score 751; DB 3; ilarity 97.4%; Pred. No. 4e-178; Conservative 0; Mismatches 20
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APPLICANT: Sharma, Sanjai
APPLICANT: Cantwell, Mark
TITLE OF INVENTION: NOVEL EX
                                              Similarity
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Best Local S:
Matches 763;
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4e-178;
20;
                                               MODULATION AND TREATMENT OF MALIGNANCIES
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CONTAINING ACCESSORY MOLECULE LIGAND GENES
                                     THEIR USE FOR IMMUNO-
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3R: 231/003
                                                                                                                                                                                                                                                                                                         .c. DOS
                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: IBM P.C. DOS
SOFTWARE: FastSeq Version 2.0
                                                                                                                   E: Lyon & Lyon 633 West Fifth Street Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                            60/132145
                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       95.9%;
nilarity 97.4%;
Conservative 0
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APPLICATION NUMBER: 60/13.
FILING DATE: 12/9/96
ATTORNEY/AGBNT INFORMATION:
NAME: Gulse, Jeffrey W.
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(213) 955-0440
                                                                                                                                                                                                                                                                                         IBM Compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 67-3510
TELEX: 67-3510
SEQUENCE CHARACTERISTICS:
ELENCTH: 783 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Guise, Jeffrey W
REGISTRATION NUMBER: 3
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NUMBER OF SEQUENCES:
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TGA 795
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Matches 763;
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 US-08-234-580-1
                                      Query Match
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ACTGAAGCAAGCCAAGTGCATCCACAGAGTTGGCTTCTCTTTTTGGCTTACTCAAACTC 780
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                                                 ACGGTTAAAAGAGAAGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGG
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                                                                                                                                    GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCT
                                                                                                                                                 APPLICANT: KEHRY, MERILYN R
APPLICANT: CASTLE, BRIAN E
TITLE OF INVENTION: METHODS FOR PROLIFERATING AND
TITLE OF INVENTION: DIFFERENTIATING B CELLS, AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEE: STERN, KESSLER, GOLDSTEIN & FOX : 100 NEW YORK AVE. N.W. SUITE 600 WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1011.1030000/RAM
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                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/08234580 GENERAL INFORMATION:
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPALIBLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: MILLMAN, ROBERT A REGISTRATION NUMBER: 36,217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION
TELEPHONE: (202)371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 28-APR-1994
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 818 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: WALL STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY:
LOCATION:
                                                                                                                                                                                                                                                                                                                                   TGA 783
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                                        Gaps
                                                                             1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
                                                                                                                    ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC 72
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                                      ö
Length 818
                                      Indels
Score 751; DB 6;
Pred. No. 4.1e-178;
                                      20;
                                      0; Mismatches
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NOVEL CYTOKINE
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GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
APPLICANT: FARMITAGE, MILLIAM
APPLICANT: FARMITAGE, MILLIAM
APPLICANT: FARMITAGE, MILLIAM
95.98;
                                        Conservative
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                      Similarity
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TCTGTTCACTTGGGCGGAGTGTTTGAALTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
                                                                                                                                                                                                                                                    ACGGITAAAAGAAGAAGGACICIATTATGICIACACICAAGICACCIICTGCICIAAICGG 540
            541 GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCT
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                                                                          421 AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCE: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/107,353
PTIING DATE: 19930813
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Immunex Corporation
51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 1003
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 5, Application US/08107353
GENERAL INFORMATION:
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nucleic acid
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INFORMATION FOR SEQ ID NO:
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HYPOTHFMITT
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ORIGINAL SOURCE:
ORGANISM: MOU
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STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 ATGATCGAAACATACAACCTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60,
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Pred. No. 1e-177;
0; Mismatches 21;
                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                             APPLICATION NUMBER: US/07/969,703A
FILING DATE: 19921023
CLASSIFICATION: 435
                                                                                                                                                          PC-DOS/MS-DOS
                  ADDRESSEE: IMMUNEX CORPORATION STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                    NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REPERNEC/POCKET NUMBER: 280;
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
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ilarity 97.3%;
Conservative
                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
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SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
TYPE: NUCLEIC ACLD
STRANDEDNESS: single
      CORRESPONDENCE ADDRESS:
                                                                                                              COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: CDNA
                                                                  WASHINGTON
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Best Local Similarity
Matches 762; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                     linear
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IMMEDIATE SOURCE:
CLONE: CD40-L
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, LOCATION: 1..7
US-07-969-703A-1
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                                                                                COUNTRY:
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THAT BINDS CD40

E: IMMUNEX CORPORATION
51 UNIVERSITY STREET

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APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIYASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
                                                                NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS:
                                                                                       ADDRESSEE:
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                                                                                                                                                         1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
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                                                                                         Length 783;
                                                                                       Score 749.4; DB 5;
Pred. No. 1e-177;
0; Mismatches 21;
                                                                                       tch 95.7%; Score 749.4; al Similarity 97.3%; Pred. No. 1e-762; Conservative 0; Mismatches
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US-08-477-733A-1
; Sequence 1, Application US/08477733A
; GENERAL INFORMATION:
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CD40-L
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                       ; NAME/KEY:
; LOCATION:
US-08-107-353-5
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STRAIN:
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Matches 76
            FEATURE
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APPLICANT: ARMITAGE, RICHARD

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61 ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
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                                                                                                       SOFTWARE: Apple Operating System 7.1 SOFTWARE: Microsoft Word for Apple, version 5.1a APPLICATION DATA: APPLICATION NUMBER: US/08/477 777 CLASCY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 749.4; DB 8;
Pred. No. 1e-177;
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                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: December 5, 1991
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783.707
FILING DAFF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            AFFLING DATE: OCTOBER: 07/783,707 FILING DATE: OCTOBER 25, 1991 CLASSIFICATION: 435 ATTORNEY-AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                UMBER: 07/969,703
October 23, 1992
                                                                                                                                                                                                                                                                08/249,189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95.7%;
97.3%;
                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
                                                                                                                                                                                                                                                                            May 24, 1994
N: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             : 783 base pairs
nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 97.3
Matches 762; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
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ADDA...
STREET: 51 ...
CITY: SEATTLE
THE: WASHINGTON
                                                                                                                                                                                                                               CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2065870606
                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER:
                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                FILING DATE: Ma
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                                                                                                                                                                                                                                                                                                                                                 FILING DATE:
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US-08-477-733A-1
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Word for Apple, version 5.1a

08/249,189

APPLICATION NUMBER: 07/805,723 FILING DATE: December 5, 1991

"CLASSIFICATION: 514 PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,70:
FILING DATE: October 23, 1992

07/783,707

October 25, 1991

TELECOMMUNICATION INFORMATION:

2065870606

TELEFAX:

MOLECULE TYPE: CDN

STRANDEDNESS:

IMMEDIATE SOURCE ANTI-SENSE: NO ORIGINAL SOURCE

CLONE

ORGANISM

NAME/KEY: LOCATION:

REGISTRATION NUMBER: 34, REFERENCE/DOCKET NUMBER:

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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/no'.-
FILING DATE:
CLASS.--
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FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATE:
APPLICATION NUMBER: 08/249,1
                                                                                                                                                                                                                                    CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,7
FILING DATE: October 25, 199
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 783 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-484-624-1
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                                                                                                                                                                             AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG 480
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241 TIGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
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                                                                                                                                  901 AACAAAGAAGAAAAAAAAAAAAAAGCTTTGAAATGCAAAGAGGTGATGAGAGTCCTCAA 360
                                                        1 ATGAICGAAACATACAACCAAACTICTCCCCGGATCIGCGGCCACTGGACTGCCCAICAGC 60
                                                                        1 ATGATAGAAACATACAGCCAAACCTTCCCCCAGATCGGTGGGAACTGGGACTTCCAGCGAGC
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                            21;
Score,749.4; DB Pred. No. 1e-177; 0; Mismatches 2
    95.7%;
                                 Matches 762; Conservative
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APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTCKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 24
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E: IMMUNEX CORPORATION 51 UNIVERSITY STREET

WASHINGTON

STATE: WALLE 98101

COUNTRY:

CORRESPONDENCE ADDRESS: ADDRESSEE: IMMUNEX C

COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: Apple Macintosh

Sequence 1, Application US/08484624 GENERAL INFORMATION:

US-08-484-624-1

APPLICANT: ARMITAGE, RICHARD

FANSLOW, WILLIAM SPRIGGS, MELANIE

APPLICANT: APPLICANT:

240

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Gaps

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Length 783;

DB 8;

PRIOR AP	FILING CLASSI PRIOR AP APPLICA	FILING CLASSII ATTORNEY, NAME:	REGIST REFERE TELECOMMI	TELEFA INFORMATION SEQUENCE LENGTH	TYPE: STRANDI TOPOLO MOLECULE	HYPOTHET ANTI-SEN ORIGINAL ORGANI	IMMEDIAT CLONE: FEATURE: NAME/K:	; US-08-770-981	Query Match Best Local	hes 76	1 ATG	1 ATG	61 ATG	61 ATG	121 CTT	121 CTT	181 GAA	181 GAA	241 TTG	241 TTG	301 AAC	301 AAC	361 ATT	361 ATT	421 AAG	421 AAG	481 ACG	481 ACG
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GGTGAT	TCCGTT TCCGTT	GAAAAT GAAAAT	ACCTTC ACCTTC	AAGCCC AAGCCC	CAGCTT CAGCTT	TCTGTG 	TTTGGC TTTGGC		,					,	CD40			. ·		•	5.1a							
CAAAGA	GCAGCA GCAGCA	ATGCTT 	CAAGTC 	TGGCTG	TCCTCC	GGTGCT GGTGCT	TCATCT 	•												P 7 1	10u	·						•
GAAATG	AGTAAT AGTAAT	TTGGTA TTGGTA	TACACT	GGCCTC GGCCTC	CACAGT CACAGT	CAAGCT 	GGCTTC								NOVEL CYTOKINE THAT BINDS 24					Svatem	~							
AGCTTT	GCCAAC GCCAAC	AGCAAC AGCAAC	TATGTC TATGTC	ATCGTC ATCGTC	AATACC AATACC	GAATTA GAATTA	AGAGTT AGAGTT				981		CUTNI	TNITH	FORTNE	TION			٠	pratin	Microsoft Word for Apple,	MBER: US/08/770,981 20-DEC-1996		,733	,189		19,703	; !
SAAAAC	AGCGAA AGCGAA	ATGAAA 	CTCTAT	CATTC	SCGGCA SCGGCA	STGTTT 	ATCCAC 	•		•	/08770	RICHARD	MELANIE	CLOU	EL CY	IMMUNEX CORPORATION			, i o i	tosh	vord for	780/st		08/477,733 1995	08/249,189		07/969,703	
AAAAAA	STTGTA	ratacci ratacci	SAAGGAG 	PAACGC 	TCAAGO TIIII	SGCGGA(CAAGTG				ton US,	ATION: ARMITAGE, RICHAR FANSIOM WILLIAM	SPRIGGS, MEI	2	й	I IMMUNEX COF	ENG!		E FORM:	Apple Macintosh	Soft N	BER: (435 DATA:	MBER: 08/4 07-JUN-1995	4	435 DATA:		435
IIIII	CACAC	GATAT	AAAGAC	CGAGT	TCTTAC TCTTAC	ACTTGC	CAAGCC				licat	ARMITAGE,	SPRIG	GIBSON	OUENCE OUENCE OF ADI	IMM	SEATTLE WASHINGTON	USA	ABI	Apple SYSTEN	Micro	N NUM	TION:	N NUM	N NUM	TION:	N NUM	TION:
	ATTGCAGCACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC 	aagaaaggatattataccatgaaagcaacttggtaatgcttgaaaatgggaaacagctg 	acgettaaaagagaagactctattatgtctacactcaagtcaccttctgctctaatcgg 	GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCT 	GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCCAGCTTTGCGAGCAGCAG 	TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAAGGTG 	ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTTATTGGCTTACTCAACTC 	TGA 783	TGA 783	181-1	1, App	ANT:		ANT	TITLE OF INVENTION: NO NUMBER OF SEQUENCES: 2	ADDRESSEE:	121 3	COUNTRY: U			SOFTWARE: Microsoft WO	APPLICATION NUMBER: FILING DATE: 20-DE	CLASSIFICATION: 435 PRIOR APPLICATION DATA:	APPLICATION NUMBER: FILING DATE: 07-JU	APPLICATION NUMBER: FILING DATE: May 2	CLASSIFICATION: 435 PRIOR APPLICATION DATA	APPLICATION NUMBER: FILING DATE: Octobe	CLASSIFICATION:
301 A	361 A	421 A 421 A	481 A	541 G	601 G - 601 G	661 T 661 T	721 A 721 A	781 T	781	SULT 10 -08-770-981-1	Sequence 1, Application US/08770981	APPLICANT:	APPLICANT:	APPLICANT:	TITLE NUMBER CORRES	ADDR	CITY: S	COUN	COMPUTER	COMP	SOFT	APPL	CLAS	APPL	APPL	CLAS PRIOR	APPI FILI	CLAS
QQ	Qy Db	Oy Dp	Qy Db	Oy Dp	Oy Dp	Qy	Oy Dp	Qy	QQ	RESULT US-08-7	Sec :	5 	· ·• ·															

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AGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATC 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  h 95.7%; Score 749.4; DB 11; Length 783; Similarity 97.3%; Pred. No. 1e-177; 62; Conservative 0; Mismatches 21; Indels 0;
PPLICATION DATA:
CATION NUMBER: 07/805;723
G DATE: December 5, 1991
IFICATION: 435
                                                            PPLICATION DATA:
CATION NUMBER: 07/783,707
G DATE: October 25, 1991
IFICATION: 435
                                                                                                                                              :: Perkins, Patricia A.
STRATION NUMBER: 34,683
RENCE/POCKET NUMBER: 2802
RENCE/TOCKET NUMBER: 2802
SPHONE: 2065870430
SPAX: 2065870606
                                                                                                                                                                                                                                                TON FOR SEG 1D NO: 1:
CE CHARACTERISTICS:
TH: 783 base pairs
:: nucleic acid
NDEDNESS: single
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NISM: MOUSE
ATE SOURCE:
E: CD40-L
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Gaps

us-08-982-272-3.rnpm

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541 GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCT 600
541 GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCAGTGGATCT 600
                                                                             TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
                                                                                                                                          721 ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC 780
                                                              GAGAGAATCTTACTCAAGGCGGCAAATACCCACACTTCCTCCCAGCTTTGCGAGCAGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SYSTEM: Apple Operating System:7.1
Microsoft Word for Apple, version 5:la
                                                                                                                                                                                                                                                                                                                                                                                                                              STATION: SUBHASHINI GIBSON, MARYLOU GIBSON, MARYLOU GIBSON: NOVEL CYTOKINE THAT BINDS CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/088,913
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PRIOR APPLICATION DATA:
APPLICATION NUMBER
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51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 07/969,70:
FILING DATE: October 23, 1992
PRIOR APPLICATION DATA:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,70
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/09088913 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PELECOMMUNICATION INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                        ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                   FANSLOW, WILLIAM SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 2 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEATTLE
WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION
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APPLICANT:
APPLICANT:
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US-09-088-913-1
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ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC 780
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                                                                                                                                                  Score 749.4; DB 14; Length 783;
Pred. No. 1e-177;
0; Mismatches 21; Indels 0;
                                                                                                                                                                          0; Mismatches
                                                                                                                                                        95.7%;
milarity 97.3%;
Conservative 0
nucleic acid
                   linear
                                                                                                                                                           Query Match
Best Local Similarity
Matches 762; Conserv
                                                                    ORGANISM: MOUSE
IMMEDIATE SOURCE:
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                                                ANTI-SENSE: NO ORIGINAL SOURCE:
          STRANDEDNESS:
                           MOLECULE TYPE: HYPOTHETICAL:
                                                                                                          ; NAME/KEY:
; LOCATION:
US-09-088-913-1
                    COPOLOGY:
                                                                                          CLONE:
                                                                                                 FEATURE:
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                                                                            ATGAAAATTTTTATGTATTTACTTACTTTTTTTTTTTATCACCAGATGATTGGGTCAGA 120
APPLICANT: ARMITMGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3: IMMUNEX CORPORATION 51 UNIVERSITY STREET
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US-09-322-021A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, version 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/322,021
                                                                                                                                                                          APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/249,189
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APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
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NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: October 25, 1991
CLASSIFICATION:
                                                                                                                Sequence 1, Application US/09322021 GENERAL INFORMATION:
                                                                                                                                                           ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
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EDNESS: single
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IMMEDIATE SOURCE:
CLONE: CD40-L
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US-09-322-021-1
                  781 TGA 783
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COUNTRY:
                                                                                                                                                         APPLICANT:
                                                                                             US-09-322-021-1
                                                                            RESULT
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WASHINGTON SEATTLE

STREET: CITY: SE STATE:

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TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
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                  421 AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAAACAGCTG
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                                                              361 ATTGCAGCACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC
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MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: MS Word for Apple Power Macintosh, Version 6.0.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: MORRIS, ARVIA E.
APPLICANT: MCGREW, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
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GIBSON, MARYLOU
MORRIS, ARVIA E.
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APPLICATION NUMBER: 07/969,703
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              October 23, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/09365940
GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FANSLOW, WILLIAM
SPRIGGS, MELANIE
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APPLICATION NUMBER:
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Pred. No. 1e-177;
0; Mismatches 21; Indels
                                                                                          OPERATING SYSTEM: Apple Operating System 7.1 SOFTWARE: Microsoft Word for Apple, version 5.1a CURRENT APPLICATION DATA:
                                                                                                                                         US/09/322,021A
                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
                                                                                                                                                                                                                                   CLASSIFICATION:
PRIOR APPLICATION APPLICATION AND APPLICATION WINBER: 07/805,723
FILING DATE: December 5, 1991.
                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/249,189
                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: PERKIAS, PALITAGIA A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802
TELECOMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEPHONE: 2065870430
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milarity 97.3%;
Conservative 0
                                                                           Apple Macintosh
SYSTEM: Apple O
                                                                   Floppy disk
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nucleic acid
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                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                           APPLICATION NUMBER:
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Matches .762; Conserv
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IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
                                                                                                                                                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; NAME/KEY: CDS
; LOCATION: 1..7
US-09-322-021A-1
                                                                                                                                                                           CLASSIFICATION:
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                                                                                                                                                                                                                        FILING DATE
                                                                                                                                                              FILING DATE
                                       98101
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                                                                                      COMPUTER:
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661 TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG
ACGGTTAAAAAGAAGAACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGG
                                                   GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCAGTGGATCT
                                                                                                                                                                                                       ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC
                            541 GAGCCTTCGAGTCAACGCCCATTCATCGTCGCCTCTGGCTGAAGCCCAGCATTGGATCT
                                                                                       GAGAGAATCTTACTCAAGGGGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/09365940A GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ouery Match
Best Local Similarity 97.33
Matches 762; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , COUNTY (1) . (7) ; OTHER INFORMATION: US-09-365-940A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: CDS
LOCATION: (1).
                                                                                                                                                                                                                                                                  781 TGA 783
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US-09-365-940A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        95.7%; Score 749.4; 97.3%; Pred. No. 1e-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
CLASSIFICATION:
                        07/805,723
                  APPLICATION NUMBER: 07/805,72
FILING DATE: December 5, 1991
                                                                                                                           ATTORNEY/ACENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERRESTICS: LENGTH: 783 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      762; Conservative
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                                                                                                                                                                                                                                                                                             single
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ORGANISM: MOUSE
IMMEDIATE SOURCE:
CLONE: CD40-L
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                                                      CLASSIFICATION:
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HYPOTHETICAL:
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; LOCATION:
US-09-365-940-1
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APPLICANT: ARMITAGE, Richard, J.
APPLICANT: FANSLOW, III, William, C.
APPLICANT: FFANSLOW, Melanie, K.
TITLE OF INVENTION: METHOD OF TREATING AN UNDESIRED IMMUNE RESPONSE BY BLOCKING TH
TITLE OF INVENTION: OF CD40 LIGAND TO CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 17; Length 783
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 749.4; DB 17;
Pred. No. 1e-177;
0; Mismatches 21;
                                                                                                                                                                                                                                                                                                                                                                                                                            FILE REFERENCE: 2802-K
CURRENT APPLICATION NUMBER: US/09/365,940A
CURRENT FILING DATE: 1999-08-02
PRIOR APPLICATION NUMBER: 08/249,189
PRIOR FILING DATE: 1994-05-24
PRIOR FILING DATE: 1992-10-73
PRIOR FILING DATE: 1992-10-23
PRIOR FILING DATE: 1992-10-23
PRIOR FILING DATE: 1992-10-65
PRIOR PALLICATION NUMBER: 07/805,723
PRIOR PILING DATE: 1991-12-05
PRIOR FILING DATE: 1991-10-25
PRIOR FILING DATE: 1991-10-25
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		241 ttgctgaactgtgaggagatgaggaggaggcaattlgaayacttgccagscarss	301 AACAAAGAAGAAAAAAAGAAAACAGCTTTGAAATGCAAAGGGGGGGG	301 aacaaagaagaagaaaaagaaaacagctttgaaatgcaaagagglugaryayyacccccc	361 ATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCGGTTCTACAGTGGGCC	361 attgcagcacacgttgtaagcgaagccaacagtaatgcagcatccgttctacagugggcu	421 AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG	421 aagaaaggatattataccatgaaaagcaacttggtaatgcttgaaaauyyyaaacuyyee	481 ACGGTTAAAAGAGAAGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGG	481 acggitaaaagagaaggactctattatgtctacactcaagtcaccttctgctctaatcg	541 GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCT	541 gagecttegagteaaegeeeatteategteggeetetggetgaageeeageagtggate	601 GAGAGAATCTTACTCAAGGGGGCAAATACCCACGTTCCTCCCAGCTTGCGAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCA	601 gagagaatettaetteaaggeggeaaataeeeacagtteeteeeagetttgegageagea	661 TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGTTCTGTGTTTGTCAACGTG	661 tctgttcacttgggcggagtgtttgaattacaagctggtgcttctgtgtttgtcaacgtg	721 ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC	721 actgaagcaagccaagtgatccacagagttggcttctcatcttttggcttactcaaactc	781 TGA 783	.781 tga 783	
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Search completed: May 30, 2002, 05:22:46 Job time: 18645 sec

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May 30, 2002, 01:00:26 ; Search time 627.19 Seconds (without alignments) 1310.273 Million cell updates/sec
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1 ATGATCGAAACATACAACCA.....TTGGCTTACTCAAACTCTGA 783
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                              OM nucleic - nucleic search, using sw model
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Sequence:
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Database :

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

length: 0 length: 2000000000

Minimum DB : Maximum DB :

Pending_Patents_NA_New:*

1: /cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*

2: /cgn2_6/ptodata/1/pna/USO6_NEW_COMB.seq:*

3: /cgn2_6/ptodata/1/pna/USO7_NEW_COMB.seq:*

4: /cgn2_6/ptodata/1/pna/USO8_NEW_COMB.seq:* /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq: /cgn2_6/ptodata/1/pna/US60_NEW_COMB.seq: Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		æ			SUMMARIES	
Result No.	t Score	Ouery Match	Length	DB	. QI	Description
	1 749.4	95.7	1250	2	US-09-053-375B-821	Sequence 821, App
	2 592.4	75.7	1816	2	US-09-053-375B-226	
	3 592.4	75.7	1816	Ŋ	US-09-442-384B-457	457,
	4 315	40.2	208	2	US-09-911-904-131	
	5 155.8	19.9	2395	Ŋ	US-09-875-453A-9	
υ	6 41.6	5.3	2865	9	US-10-027-632-111808	
U	7 39.2	5.0	1223197	9	US-10-027-632-179264	a)
υ	8 38.4	4.9	45698	9	US-10-105-299-12515.	Sequence 12515, A
υ	9 38.4	4.9	99599	'n	US-09-830-706B-37	Sequence 37, Appl
7	0 38	4.9	476	9	US-10-027-632-187249	
1	38	4.9		9	US-10-027-632-187250	18725
o I	2 38	4.9	89	9	US-10-027-632-35082	Sequence 35082, A
1	3 . 37.6	4.8	349	Ś	US-09-721-544-11180	
-	4 37	4.7	1758	9	US-10-071-192-29	
c 1	5 36.6	4.7	341	Ŋ	US-09-789-189-555	555,
-		4.7	57649	ø	US-10-105-299-9151	Sequence 9151, Ap
7		4.7	57649	9	US-10-105-299-10646	10646
1	18 36.6	4.7	57649	9	US-10-105-299-14:274	Sequence 14274, A
1		4.6	2430	9	US-10-105-694-1	Sequence 1, Appli
~		4.6	2430	9	US-10-105-695-1	٦,
2	1 36.2	4.6	2430	9	US-10-106-014-1	ī
7	36.	4.6	3321	Ŋ	US-09-919-002-413	413
•	3	4.6	1379	9	US-10-027-632-199307	
0	35.8	4.6	598	9	US-10-027-632-207524	Sequence 207524,
0	5 35.8	4.6	2580	9	US-10-027-632-258659	Sequence 258659,
c 2	9.32.6	4.5	626	9	US-10-027-632-239684	Sequence 239684,

	Sequence 1404, Ap	Sequence 499, App	Sequence 8655, Ap	Sequence 219364,	Sequence 219365,	Sequence 233265,	Sequence 141134,	Sequence 380, App	Sequence 270, App	Sequence 362, App	Sequence 362, App	Sequence 85186, A	Sequence 85187, A	Sequence 46525, A	Sequence 46526, A	Sequence 210428,	Sequence 200394,	Sequence 228875,	Sequence 233425,
	US-10-106-698-1404	US-10-106-698-499	US-09-919-002-8655	US-10-027-632-219364	US-10-027-632-219365	US-10-027-632-233265	US-10-027-632-141134	US-10-113-872-380	US-09-053-375B-270	PCT-US02-07826-362	US-10-097-340-362	US-10-027-632-85186	US-10-027-632-85187	US-10-027-632-46525	US-10-027-632-46526	US-10-027-632-210428	. US-10-027-632-200394	US-10-027-632-228875	US-10-027-632-233425
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	712	1640	1665	604	604	637	856	522	3304	3310	3310	601	601	620	620	1216	584	601	608
	4.5	4.5	4.5	4.5	4.5	4.5	4.5	4.4	4.4	4.4	4.4	4.4	4.4	4.4	4.4	4.4	4.4	4.4	4.4
	35.6	35.6	35.6	35.2	35.2	35.2	35	34.8	34.8	34.8	34.8	34.6	34.6	34.6	34.6	34.6	34.4	34.4	34.4
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ALIGNMENTS

Query Match 95.7%; Score 749. Best Local Similarity 97.3%; Pred. No. Matches 762; Conservative 0; Mismatc	Oy 1 ATGATCGAAAGATACAACCAAACTTCTCCCCGATC	Oy 61 ATGAAATTTTATGTTTACTTACTGTTTTCTTATCACCCAGATGATTGGGTCAGCA 120	Oy 121 CTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAGGAAGACTTCAT
. 10	 	Juery March Best Local Similarity Matches 762; Conservat I ATGATCGAAACATACA 	Matches Aatches 1 13 73

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GAAGATTTTGTATTCATAAAAAAGCTAAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC

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ATTGCAGCACATGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC 420

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APPLICANT:
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                                                      tetgtteacttggggggggggtgtttgaattacaagetggtgettetgtgtttgteaaegtg
                                                                                                      GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCT
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Pred. No. 1.8e-142;
0; Mismatches 111;
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CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 1543
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 226
                                                                                                                                                                                                                                                                                                                            APPLICANT: Chenchik, Alex
APPLICANT: Bibilashvilli, Robert
TITLE OF INVENTION: Nucleic Acid Arrays
FILE REFERENCE: CLON-006
                                                                                                                                                                                                                                                                                                             Sequence 226, Application US/09053375B
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85.5%;
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US-09-053-375B-226
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US-09-053-375B-226
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Best Local S:
Matches 672,
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                                                                                         GCCAAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAG
                        CIGACGGITAAAAGAAGAAGGACICIATIATGICIACACCICAAGICACCITCIGCICIAAI
AACAAAGAAGA---GAAAAAAGAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCT
                                                                    CAAATTGCAGCACGCTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGG
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1.8e-142;
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CURRENT APPLICATION UNMER: US/09/442,384B
CURRENT FILING DATE: 1999-11-17
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Pred No 1.8e-
0; Mismatches
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SOFTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Chenchik, Alex
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PRIOR APPLICATION NUMBER: 09/053,375
PRIOR FILING DATE: 1998-03-31
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85.5%;
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Best Local Similarity 85.5
Matches 672; Conservative
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; ORGANISM: homo sapiens
US-09-442-384B-457
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                                    GAAGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
                                                                       TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA
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Pred. No. 1.9e-71;
0; Mismatches 120; Indels
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APPLICANT: Neft, Robin Bileen
APPLICANT: Dunn, II, Robert Thomas
TILLE OF INVENTION: CANINE TOXICITY GENES
FILE REFERENCE: 400742000200
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PRIOR FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 386
SOFTWARE: FastSEQ for Windows Version 4.0
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Best Local Similarity 76.3%;
Matches 387; Conservative
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US-09-911-904-131
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LENGTH: 508
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2 caatttgaagcctttctcaaggagataatgctaaacgaaacgaaatgaagaagaagaaaac
                                                   328 TTTGAAATGCAAAGAGGTGATGAGGATCCTCAAATTGCAGCACGCGTGTAAGCGAAGCC
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TITLE OF INVENTION: PROMOTERS FOR REGULATED GENE EXPRESSION
FILE REPERENCE: 54600-8135.US00
CURRENT APPLICATION UNMBER: 2009-06-06
PRIOR APPLICATION NUMBER: US 60/209,549
PRIOR FILING DATE: 2000-06-06
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Pred. No. 2.3e-30;
0; Mismatches 17;
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90.7%;
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Sheppard, Liana 1
Lim, Moon Young
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Thomas, Rita L.
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Best Local Similarity 90.7
Matches 166; Conservative
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US-09-875-453A-9
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Db 759231 TCCGTTGCAATGSGGAGTGACGTTTATCCCTTCYTRATAGGGAGGCAAATATTTGGGTCA 759172
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 212 GATGCAACAAAGGAAGGATCTTTATCCTTGCTGAACTGTGAGGAGATGAGAAGGCAAT 271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92 ITCTTATCACCCAGATGATTGGGTCAGCACTTTTTGCTGTGTATCTTCATAGAAGATTGG 151
                                                                             APPLICANT: Wang, David G. Titleation and Mapping of Single Nucleotide TITLE OF INVENTION: Identification in the Human Genome TITLE OF INVENTION: Polymorphisms in the Human Genome FILE REFERENCE: 108827.129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 39.2; DB 6;
Pred. No. 14;
3; Mismatches 106;
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TITLE OF INVENTION: Human Secreted Proteins
FILE REFERENCE: PS950
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 272 TIGAAGACCTIGICAAGGAIATAACGITAA 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                               CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR APPLICATION NUMBER: US 60/218,006
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                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 1999-09-28
APPLICATION NUMBER: US 60/146,002
FILING DATE: 1999-08-09
                                                                                                                                                                                                                               FILING DATE: 2000-07-12
APPLICATION NUMBER: US 60/198,676
                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US 60/185,218
                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 2000-02-24
APPLICATION NUMBER: US 60/167,363
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                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 60/193,483
                                       , Application US/10027632
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US-10-027-632-179264
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Matches 101; Conservative
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SEQ ID NO 179264
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                                                                                    0; Gaps
                                                           ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCÁGATGATTGGGTCAGCA 120
                                                                                                                                           121 CTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAAGAAGTAAACCTTCAT 180
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TITLE OF INVENTION: Identification and Mapping of Singititle OF INVENTION: Polymorphisms in the Human Genome FILE REFERENCE: 108027,129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
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Pred. No. 0.5
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FastSEQ for Windows Version 4.0
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IOR FILING DATE: 2000-07-12

IOR PILING DATE: 2000-07-13
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APPLICATION NUMBER: US 60/146,002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 60/198,676 FILING DATE: 2000-04-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US 60/193,483
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ilarity 46.8%;
Conservative
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Matches 131; Conserv
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US-10-027-632-111808
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Length 1223197;

us-08-982-272-3.rnpn

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TITLE OF INVENTION: Ide
TITLE OF INVENTION: PC
FILE REFERENCE: 108827
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                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Human
US-10-027-632-187249
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                                                                                                                                                                                    217 AACAAAGGAGGAACCTTTATCCTTGCTGAACTGTGAGGAGGAAGGCAATTTGAA 276
                                                                                                                                                                                                                                                          248 ACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAG 307
                                                                                                                                                    Gaps
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                                                                                                                Length 45698;
                                                                                                          Score 38.4; DB 6; Length 4 Pred. No. 8; 0; Mismatches 91; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: YANO, MINORU
APPLICANT: TAMAI, KATSUYUKI
TITLE OF INVENTION: THIOREDOXIN REDUCTASE II
FILE REFERENCE: 55865-71965
CURRENT APPLICATION NUMBER: US/09/830,706B
CURRENT FILING DATE: 2002-01-29
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Pred. No. 9;
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GENERAL INFORMATION:
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; OTHER INFORMATION: n is unknown
US-09-830-7068-37
                                                                                                            4.9%;
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SOFTWARE; Patentin Ver. 2.1
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                                                                                                                                                93; Conservative
                   ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-105-299-12515
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                                                                                                            Query Match
Best Local Similarity
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SOFTWARE; Patent.
SEQ ID NO 37
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LENGTH: 45698
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Matches 6
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GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILER REFERENCE: 108827.129
CURRENT APPLICATION WHERE: US/10/027,632
CURRENT FILING DATE: 2002-04-30
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Pred. No. 2.4;
0; Mismatches 100;
                                             PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1909-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,363
PRIOR FILING DATE: 1999-11-23
PRIOR PLING DATE: 1999-11-3
PRIOR PLING DATE: 1999-10-28
PRIOR FILING DATE: 1999-10-29
PRIOR FILING DATE: 1999-10-29
CURRENT APPLICATION NUMBER: US/10/027,632 CURRENT FILING DATE: 2002-04-30
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PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
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PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR PRILING DATE: 2000-02-24
PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
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Best Local Similarity 49.5
Matches 98; Conservative
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389 ACAGTAATGCAGCATCCGTTCTACAGTGGGCCAAGAAAGGATATTATACCATGAAAAGCA 448
                                                                                                                                                            318 AAAATGAAAAATCATGTTAAAAACTTGTGCAAAAGTGTTGATATTAAAATATAAAATACTAA 259
                                       329 TIGAAATGCAAAGAGGTGATGAGGATCCTCAAATTGCAGCACGTTGTAAGCGAAGCCA 388
438 AAAGTGCCTACTTTATCCTAGATATCATGTGAGGCAATGTTGAAAATATGATGAAAAGAA 379
                                                                             378 CAGATTTTCAATGACATAATAAGGCTTATCATAGAGCAGAAGAAAGCTGAAATACACAA
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                                                                                                                                                                                                                                                                                                               Single Nucleotide
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Pred. No. 2.7;
0; Mismatches 100; Indels
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TITLE OF INVENTION: Identification and Mapping of Sing
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
                                                                                                                                                                                                               Score 38; DB 6; Le
pred. No: 2.4;
0; Mismatches 100;
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JMBER: US 60/185,218
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APPLICATION NUMBER: US 60/146,002
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US 60/167,363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US 60/198,676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  , Application US/10027632
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             aatggagaataataaaa 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            449 ACTTGGTAATGCTTGAAA 466
                                                                                                                                                                                                                     Query Match 4.9%;
Best Local Similarity 49.5%;
Matches 98; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DATE: 2000-02-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ), DOCALLON: (1)...(003)
), OTHER INFORMATION: D
US-10-027-632-35082
                                 NUMBER OF SEQ ID NOS:
SOFTWARE: FastSEQ for
SEQ ID NO 187250
                                                                                                                   ; TYPE: DNA
; ORGANISM: Human
US-10-027-632-187250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 35082
LENGTH: 685
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                                                                                                  LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      219
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, Servando ariba

Margie

Benjamin

Michelle

80, Application US/09721544

Damavandi, Simin Dickson, Mark Drake, Jim

rmanac, Radoje

Brandy

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256 GAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAGAAGAAA 315
                                                                                                                                                                                                                                                                                                                                                                                           316 AAAGAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAAATTGCAGCACACGTT 375
                                                                                                                                                                                                                                                                                                      Length 349;
lovel Nucleic Acid Sequences Obtained
From a CDNA Library of Fetal Liver-Spleen
                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                          DB 5;
                                                                                                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                            SEQ ID NOS: 24489
FastSEQ for Windows Version 3.0
                                                  APPLICATION NUMBER: US/09/721,544
                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 56.55
Matches, 70; Conservative
                                                                                        PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
US-09-721-544-11180
                                                                                                                                                                     SOFTWARE: FascasEQ ID NO 11180
                                                                                                                                                                                                                                                                                                                                                                                                                        qq
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Best Local Similarity 49.59 Matches 98; Conservative

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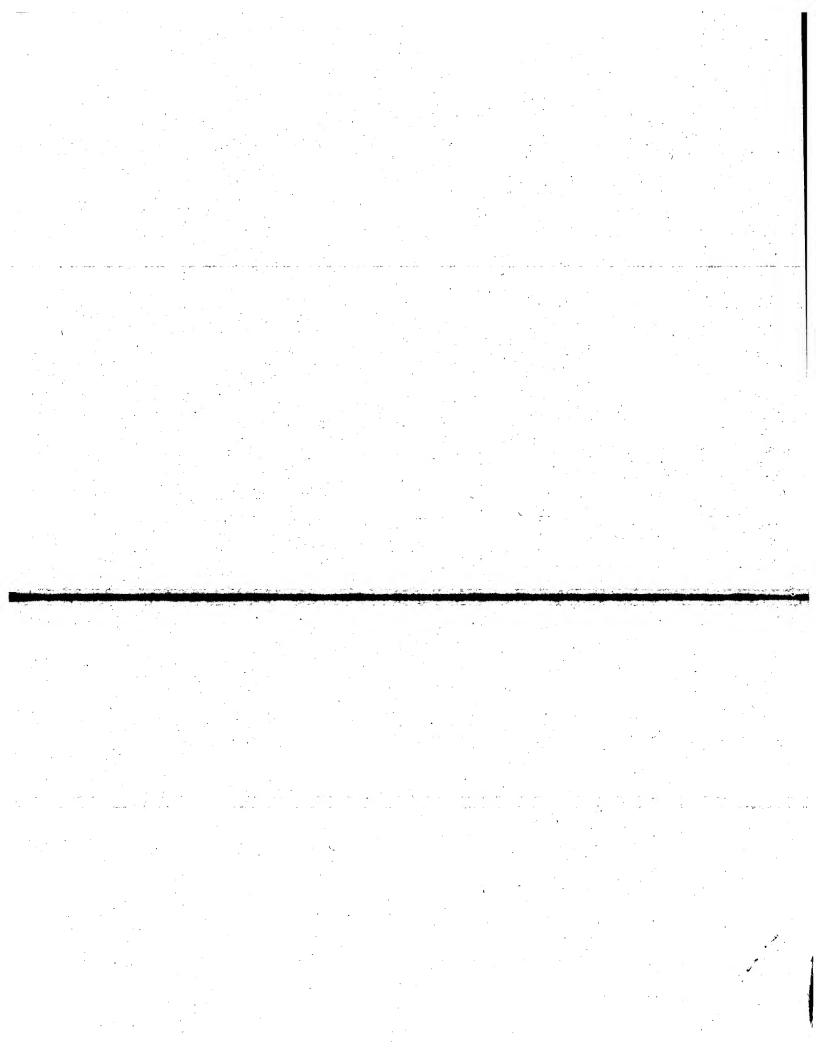
Gaps

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ORGANISM: Homo sapiens
US-09-789-189-555
                                                                                                                                                                                                                                                                                                                                                                                                                   SEQ ID NO 555
LENGTH: 341
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
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                                                                                                                                                                                                                                                        Weber, Eric R.
TITLE OF INVENTION: NOVEL ECTOPARASITE SALIVA PROTEINS AND APPARATUS TO COLLECT SUCH PROTEINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 37; DB 6; Length 1758;
Pred. No. 6.6;
1; Mismatches 121; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Connell, Gary J.
REGISTRATION NUMBER: 32,020
REFERENCE/DOCKET NUMBER: 2618-17-C4-PUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/071,192
FILING DATE: 07-Feb-2002
CLASSIFICATION: <UNKNOWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: SHERIDAN ROSS P.C.
STREET: 1560 BROADWAY, SUITE 1200
CITY: DENVER
STATE: CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; EQUENCE DESCRIPTION: SEQ ID NO: 29: US-10-071-192-29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 09/171,156
FILING DATE: 1998-10-09
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: TELEPHONE: 303/863-9700
                                                                                                                                                                            Sequence 29, Application US/10071192 GENERAL INFORMATION:
                                                                                                                                                                                                                   APPLICANT: Hunter, Shirley Wu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 1758 base pairs
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Best Local Similarity 47.2%;
Matches 109; Conservative
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                                                                                                                                                                                                                                          Sim, Gek-Kee
                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A. ZIP: 80202
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                                                376 GTAA 379
                                                                                  253 gaaa 256
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US-10-071-192-29
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                                                                                                       335 TGCAAAGAGGTGATGAGGATCCTCAAATTGCAGCACACGTTGTAAGCGAAG 385
                                                                                                                                                                                                          US-09-789-189-555/c

Sequence 555, Application US/09789189
GENERAL INFORMATION:
TITLE OF INVENTION: Human Polynucleotides and Polypeptides
FILE REPERBERS.
CURRENT APPLICATION: NUMBER: US/09/789,189
CURRENT FILING DATE: 2001-02-20
PRIOR APPLICATION NUMBER: 60/183452
PRIOR APPLICATION NUMBER: 60/183452
PRIOR FILING DATE: 2000-02-18
NUMBER OF SEQ ID NOS: 2005
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 341;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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Pred. No. 5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          158 AGGTTGTGAAAAATAAAAGATGTAATG 132
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Best Local Similarity 53.13
Matches 78; Conservative
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Sequence Sequence Sequence

Sequence Sequence Sequence

Sequence Sequence Sequence Sequence

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Appl Appl Appl Appl

Sequence Sequence Sequence Sequence

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Sequence 1, Application US/08431055
Patent No. 5817516
GENERAL INFORMATION
APPLICANT: CASTLE, BRIAN E
TITLE OF INVENTION: METHODS FOR PROLIFERATING AND
TITLE OF INVENTION: DIFFERENTIATING B CELLS, AND USES THEREOF
NUMBER OF SEQUENCES 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: STERN, KESSLER, GOLDSTEIN & FOX
STREET: 100 NEW YORK AVE. N.W. SUITE 600
STATE: D.C.
STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/431,055
FILING DATE: 28-APR-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/234,580
FILING DATE: 28-APR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 751; DB 1; 1
Pred. No. 1.3e-212;
US-09-088-913A-11

US-08-589-771B-7

US-08-769-819-11

US-08-770-974-11

US-08-858-197-3

US-08-249-189-15

US-08-477-733B-15

US-09-088-913A-15

US-08-770-974-15

US-08-770-974-15

US-08-740-15

US-08-770-974-15

US-08-448-624A-20

US-08-484-624A-20

US-08-484-624A-20

US-08-481-624A-20

US-08-481-624A-20

US-08-481-624A-20

US-08-819-20

US-08-819-20

US-08-819-20

US-08-913A-20

US-08-913A-20

US-08-913A-20
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REFERENCE/DOCKET NUMBER: 1011.1030000/RAM
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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97.48;
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TELEFAX: (202)371-2540
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: MILLMAN, ROBERT A REGISTRATION NUMBER: 36,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 818 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                           1425
1425
1425
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EDNESS: both
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Best Local Similarity
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                                                                          RESULT
                                                                                                     May 29, 2002, 23:38:16 ; Search time 139.75 Seconds (without alignments) 1376.251 Million cell updates/sec
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                                                                                                                                                                    US-08-982-272-3
783
1 ATGATCGAAACATACAACCA.....TTGGCTTACTCAAACTGGA 783
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/cgn2_6/ptodata/2/ina/6B_COMB.seq:*
/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
/cgn2_6/ptodata/2/ina/Packfiles1.seq:*
              GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /cgn2_6/ptodata/2/1na/5A_COMB.seq:*/cgn2_6/ptodata/2/1na/5B_COMB.seq:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -08-484-624A-22
                                                                                                                                                                                                                                                                                                             hits satisfying chosen parameters:
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US-08-770-974-22
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US-08-770-974-1
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                                                                                                                                                                                                                                                                                                                                                                                        Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                          nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 2000000000
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                                                                                                                                                                      Title:
Perfect score:
Sequence:
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Length 818;

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                              AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG
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  Indels
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STREET: 100 NEW YORK AVE. N.W. SUITE 600
  Mismatches
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APPLICANT: KEHRY, MERILYN R
APPLICANT: CASTLE, BRIAN E
    Conservative
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TITLE OF INVENTION:
TITLE OF INVENTION:
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                                                  Version #1.25
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                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 1011.1030000/RAM TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                    Score 751;
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                                                                                                                           APPLICATION NUMBER: US/08/234,580 FILING DATE: 28-APR-1994 ATTORNEY/AGENT INFORMATION:
                                         SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/858 107
                                      PC-DOS/MS-DOS
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                    NAME: MILLMAN, ROBERT A REGISTRATION NUMBER: 36,217
                                                                                                                                                                                                                                                                                                                                                                                                                   95.98;
                                                                                                                                                                                                                         (202)371-2600
                                                                                                                                                                                                                                  TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO:
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763; Conservative
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nucleic acid
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                                        SYSTEM:
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                                                                                                      CLASSIFICATION:
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US-08-858-197-1
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                             COMPUTER:
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                                                              601 GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
                                                                                                                                                                                                                                                                                                            Sequence 5, Application US/08446922
Patent No. 5716805
GENERAL INFORMATION:
APPLICANT: Spriggs, Melanle
APPLICANT: Strinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
TITLE OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SYSTEM: Apple Operating System 7.1 Microsoft Word for Apple, Version
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446,922
                                                                                                                                                                                                                                                                                                                                                                                                                            E: Immunex Corporation
51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 1003
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206,587-0430
TELEFAX: (206,233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: USS:
FILING DATE: 08-13-93
CLASSIFICATION: 435
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LENGTH: 783 base pairs
TYPE: nucleic acid
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STREET: 51 Univ
CITY: Seattle
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                                                                           ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
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 Length 783;
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Score 749.4; DB 1;
Pred. No. 3.9e-212;
0; Mismatches 21;
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95.7%;
97.3%;
                                      Conservative
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Patent No. 5961974
GENERAL INFORMATION:
                 Similarity
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TGA 783
Query Match
Best Local Simi
Matches 762;
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721 ACTGAAGCAAGCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAACTC 780
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                                                                                                                                          361 ATIGCAGCACACGITGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGGCC 420
                                                                                                                                                                                                   421 AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG 480
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                TTGCTGAACTGTGAGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
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                                                                                                                           181 GAAGATTTTGTATTCAFAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
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Macintosh, version 6.0.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: Apple Operating SOFTWARE: MS Word for Apple Power
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SHINIVASAN, SUBHASHINI APPLICANT: GIBSON, MARYLOU GIBSON, MARYLOU APPLICANT: MOREN, ARVIA E. APPLICANT: MORREW, JEFFERY TITLE OF INVENTION: NOVEL CYTOKIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5: IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application US/08484624A Patent No. 5962406
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Apple Macintosh
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MEDIUM TYPE: Floppy disk
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WASHINGTON
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781 TGA 783
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Pred. No. 3.9e-212;
0; Mismatches 21;
                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/249,189
          APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                                                                     FILING DATE: October 23, 1992
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
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                                                                                                                                                                                                                                                                                                                                         07/969,703
                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
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APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
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CLASSIFICATION: 514
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NAME: Perkins, Particia A.
REGISTRATION NUMBER: 34.63
REFERENCE/DOCKET NUMBER: 28(
TELECOMMUNICATION INFORMATION:
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Best Local Similarity .97.3%;
Matches 762; Conservative 0
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INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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TYPE: nucleic acid
STRANDEDNESS: single
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MEDIUM TYPE: Floppy
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IMMEDIATE SOURCE:
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ORIGINAL SOURCE:
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US-08-249-189-1
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SOFTWARE: MS Word for Apple Power Macintosh, version 6.0.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,733B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELNIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
APPLICANT: MCRRIS, ARVIA E.
APPLICANT: MCGREW, JEFFRRY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/08477733B Patent No. 5981724
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51 UNIVERSITY STREET
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October 23, 1992
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N: 435
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MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
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APPLICATION NUMBER: (
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CITY: SEATTLE
STATE: WASHINGTON
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US-08-477-733B-1
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Pred. No. 3.9e-212;
0; Mismatches 21;
APPLICATION NUMBER: US/08/484,624A
                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                   07/969,703
                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 07/783,707 FILING DATE: October 25, 1991
                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patitaia A.
REGISTRATION NUMBER: 34,693
REFERENCE/POCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEFAN: 2065870430
TELEFAN: 2065870606
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                   APPLICATION NUMBER: 07/969,70
FILING DATE: October 23, 1992
                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,
FILING DATE: June 07, 1995
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Best Local Similarity 97.3%;
Matches 762; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 783 base pairs TYPE: nucleic acid
                                                                                                                   PRIOR APPLICATION DATA:
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IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                 CLASSIFICATION:
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; LOCATION:
US-08-484-624A-1
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MEDIUM TYPE: Floppy disk
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APPLICATION NUMBER: US
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 749.4; DB 2;
Pred. No. 3.9e-212;
0; Mismatches 21;
FILING DATE.
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
TITING DATE: October 25, 1991
TITING DATE: A35
                                                                                            ATTORNEY/AGENT INFORMATION:
NAME: PERKINS, PARTICIS A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEPHONE: 2065870430
                                                                                                                                                                             TELEFAX: 2065870000
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ouery Match 95.7%;
Best Local Similarity 97.3%;
Matches 762; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                  1..783
                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                       ORIGINAL SOURCE:
ORGANISM: MOUSE
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                            ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                              HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                  ; NAME/KEY:
; LOCATION:
US-08-477-733B-1
                                                                                                                                                                                                                                                                                                                                                                                        CLONE:
                                                                                                                                                                                                                                                                                                                                                                                                        FEATURE
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661 TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG
                                                      GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: Apple Macintosh OPERATING SYSTEM: Apple Operating System 7.5.5 SOFTWARE: MS Word for Apple Power Macintosh, version
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: GIBSON, MARKLOU
APPLICANT: MORRIS, ARVIA E.
APPLICANT: MCGREW, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US/09/088,913A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SRINIVASAN, SUBHASHINI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07/969,703
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/09088913A Patent No. 6087329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         : IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA: APPLICATION NUMBER: 08/484,624
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     October 23, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              December 5, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: October 25, 1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 28
RELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 08/477, FILING DATE: June 07, 1995 PRIOR APPLICATION DATA: APPLICATION NUMBER: 07/969,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Perkins, Patrícia A
REGISTRATION NUMBER: 34,69
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us-08-982-272-3.rni

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CITY: SEATTLE
STATE: WASHINGTON
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                                                                                                                         GENERAL INFORMATION:
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CLONE: CD40-L
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                                                                                                                                                                                                                                                                                                                     USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE:
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                 781 TGA 783
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; LOCATION:
US-08-769-819-1
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STREET: 5
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APPLICANT:
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                                                                              US-08-769-819-1
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                                                                                                                                                                                                                                        Score 749.4; DB 3; Length 783;
Pred. No. 3.9e-212;
0; Mismatches 21; Indels 0
                                                                                                                                                                                                                                        Query Match 95.7%;
Best Local Similarity 97.3%;
Matches 762; Conservative
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                             TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
                                                                                                                                                                                 1..783
                                                                                                     ORGANISM: MOUSE IMMEDIATE SOURCE: CLONE: CD40-L
                                                                                                                                                                  ; NAME/KEY: CDS
; LOCATION: 1..7
US-09-088-913A-1
                                                                          ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                    FEATURE:
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APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
CORRESPONDENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Microsoft Word for Apple, version 5.1a SOFTWARE: Microsoft Word for Apple, version 5.1a Application DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/769,819
FILING DATE: 19-DEC-1996
CLASSIFICATION: 424
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FILING DATE: 07-JUN-1995
APPLICATION NUMBER: 08/249,189
FILING DATE: May 24, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: OCTOBER 23, 1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
                                                                                                                                                                                                                                                                                                                                                                                                                        E: IMMUNEX CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: 07/783,707
FILING DATE: OCLOBER 25, 1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                               Sequence 1, Application US/08769819
Patent No. 6264951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Op
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 2065870606
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 783 base pairs
TYPE: nucleic acid
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781 TGA 783
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TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            aacaaagaagaaaaaaaaaaaaaagaagctttgaaatgcaaagaggtgatgaggatcctcaa 360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GAAGATTTTGTATTCATAAAAAAGCTAAAGAGATGCAACAAAGAGAGAAGGATCTTTATCC 240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  240
                                                                                                                                                                                                                                                                                                                                       AIGAAAATITITATGTATITACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
                                                                                                                                                                                                                                 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
                                                                                                                                   Gaps
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0
Length.783;
                                                      Indels
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  Score 749.4; DB 4;
Pred. No. 3.9e-212;
0; Mismatches 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SRINIVASAN, SUBHASHINI
GIBSON, MARYLOU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Application US/08770974
Patent No. 6290972
GENERAL INFORMATION:
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        Query Match
Best Local Similarity 97.3%;
Matches 762; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ARMITAGE,
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TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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TGA 783
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Pred. No. 3.9e-212;
0; Mismatches 21;
                                                                                                                         version 5.1a
                                                                                                       Apple Operating System 7.1
                                                                                                                         Microsoft Word for Apple,
                                                                                                                                             20-DEC-1996
                                                                                                                                                                                                                      FILING DATE: 02-A0G-1995
APPLICATION NUMBER: 08/249,189
FILING DATE: May 24, 1994
PRIOR APPLICATION DATE: MAY 24, 1994
APPLICATION NUMBER: 07/969,703
                                                                                                                                                                                     CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,733
                                                                                                                                                                                                                                                                                   UMBER: 07/969,703
October 23, 1992
                                                                                                                                                                                                                                                                                                                          07/805,723
                                                                                                                                                                                                                                                                                                                                                                  07/783,707
                                                                                                                                                                                                                                                                                                                                     FILING DATE: December 5, 1991
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                            34,693
                                                                                                                                                                                                                                                                                                                                                                                                          NAME: PERKINS, PATRICIA A. REGISTRATION NUMBER: 34,69: REFERENCE/DOCKET NUMBER: 28 TÉLECOMMUNICATION INFORMATION:
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Best Local Similarity 97.3%;
Matches 762; Conservative '
                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            783 base pairs
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                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
51 UNIVERSITY
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APPLICATION NUMBER:
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APPLICATION NUMBER:
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                            WASHINGTON
                                                                                                                OPERATING SYSTEM:
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IMMEDIATE SOURCE:
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US-08-770-974-1
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                                TTGCTGAACTGTGAGGAGTGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA
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                                                                                                                                                                                                                                                                                        Sequence 5, Application PC/TUS9310034
GENERAL INFORMATION:
APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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STREET: 51 University Street
CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 1003
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Pred. No. 3.9e-212;
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                                                                                                                                                                                                                                                                                Score 749.4;
                                                                                                                                                                                                                                                                              95.7%;
97.3%;
                                                                                                  MOLECULE TYPE: CDNA to mRNA
             INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
TYPE: nucleic acid
(206)233-0644
                                                                                                                                                                                                                                                                                                            Conservative
                                                                          single
                                                                                                                              ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: MOUSE
STRAIN: CD40-L
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Matches 762; Conser
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                                                          TYPE: nucleic STRANDEDNESS:
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PCT-US93-10034-5
TELEFAX:
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                                                                                                                                       211 AGAIGCAACAAAGAAGGAACTITATCCTIGCTGAACTGTGAGGAGATGAGAAGGCAA 270
                                                                                                                                                     151 GATAAGGTCGAAGAGGAAGTAAACCTTCATGAAGATTTTGTATTCATAAAAAAGCTAAAG 210
                                                                                                     511 TACACTCAAGTCACCTTCTGCTCTAATCGGGAGCCTTCGAGTCAACGCCCATTCATCGTC
                                                                                                                                                                                                                                        331 GAAATGCAAAGAGGTGATGAGGATCCTCAAATTGCAGCACGCGTTGTAAGCGAAGCCAAC
                                                                ;
0
                                      Length 878;
                                                                   Indels
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APPLICANT: MCGREW, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
                                       Score 631.4; DB 2;
Pred. No. 3.1e-177;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   751 GGCTTCTCATCTTTGGCTTACTCAAACTGTGA 783
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Patent No. 5962406
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GIBSON, MARYLOU
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51 UNIVERSITY STREET
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                                              80.6%;
                                                                        ,Conservative
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93..857
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                                                             Similarity
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US-08-484-624A-22
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APPLICANT:
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APPLICANT:
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STATE:
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US-08-249-189-22
                                                              Best Local Sim
Matches 632;
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                                                  Query Match
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     721 ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC 780
                 SOFTWARE: Microsoft Word for Apple, version 5.1a CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: Apple Macintosh OPERATING SYSTEM 7.1
                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/249,189
                                                                                                                                                                                                                          APPLICANT: SRINTYASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
CLASSIFICATION: 51
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .07/969,703
                                                                                                                                                                                                                                                                                        5: IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: October 25, 1991
                                                                                                                                                Sequence 22, Application US/08249189
Patent No. 5961974
GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION UNBER: 07/969,70:
FILING DATE: October 23, 1992
CLASSIFICATION: 514
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LECOMMINITATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                      FANSLOW, WILLIAM
SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 878 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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15..92
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                                                                                                                                                                                                                                                                                                                                               USA
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NAME/KEY:
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LOCATION:
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                                                                   781 TGA 783
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                                                                                                                                          US-08-249-189-22
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511 TACACTCAAGTCACCTTCTGCTCTAATCGGGAGCCTTCGAGTCAACGCCCATTCATCGTC 570
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APPLICATION NUMBER: US/08/477,733B FILING DATE: June 07, 1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
APPLICANT: MORRIS, ARVIA E.
APPLICANT: MCGREW, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                             825 GCTTCTCATCTTTTGGCTTACTCAAACTCTGA 857
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APPLICATION NUMBER: 08/249,189
FILING DATE: May 24, 1994
CLASSIFICATION: 435
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APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
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APPLICATION NUMBER: 07/969,703
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
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PRIOR APPLICATION DATA
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                                               OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: MS Word for Apple Power Macintosh, version 6.0.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,624A
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Pred. No. 3.1e-177;
0; Mismatches 1;
                                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
CLASSIFICATION:
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APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
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APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
                                                                                                                                                                               08/477,733
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NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802-
TELECOMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAS: 2065870430
TELEFAS: 2065870606
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 878 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Murine CD40-L trimer
                                                                                                                                                                         APPLICATION NUMBER: 08/477, FILING DATE: June 07, 1995 CLASSIFICATION:
                 MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Op
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99.8%;
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Best Local Similarity 99.89
Matches 632; Conservative
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93..857
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LOCATION: 15..92
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PRIOR APPLICATION DATA:
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COMPUTER READABLE FORM:
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; LOCATION:
US-08-484-624A-22
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                                                                                US-09-088-913A-22
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APPLICANT:
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                                                                        RESULT
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Pred. No. 3.1e-1-77;
0; Mismatches 1;
APPLICATION NUMBER: 07/783,70°
FILING DATE: OCtober 25, 1991
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFRENCE/COCKET NUMBER: 2802
TELEPHONE: 2055870430
             October 25, 1991
                                                                                                                                                                                                                                                                                                                                                       80.6%;
nilarity 99.8%;
Conservative 0
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                                                                                                                  22:
                                                                                                      INFORMATION FOR SEO ID NO: 2% SEQUENCE CHARACTERISTICS: LENGTH: 878 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                    mat_peptide
93..857
                                                                                                                                                                                                                                 NAME/KEY: sig_peptide LOCATION: 15..92
                                                                                                                                                                  single
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Best Local Similarity
Matches 632; Conserv
                                                                                                                                                                            linear
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US-08-477-733B-22
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FEATURE:
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                                                                                                                                                                                                                CLONE:
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765 CAAGCIGGIGCITCIGIGITIGICAACGIGACTGAAGCAAGCCAAGIGAICCACAGAGIT 824
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                                                                                                                                                                                                                                                                                                                                      NOVEL CYTOKINE THAT BINDS CD40
                                       751 GGCTTCTCATCTTTTGGCTTACTCAAACTCTGA 783
                                                               825 GGCTTCTCATCTTTTGGCTTACTCAAACTCTGA
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                                                                                                                                                           Sequence 22, Application US/09088913A Patent No. 6087329
                                                                                                                                                                                                                                                                        SRINIVASAN, SUBHASHINI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08/484,624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: June 07, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,705
FILING DATE: October 23, 1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: December 5, 1991
PRIOR APPLICATION DATA:
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ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693 "REFERENCE/DOCKET NUMBER: 2802
                                                                                                                                                                                                                                                                                                                                                                                                IMMUNEX CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLONE: · Murine CD40-L trimer
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MEDIUM TYPE: Floppy disk
COMPUTER: APPIE MAGIATION
COMPUTER: APPIE MAGIATION
OPERATING SYSTEM: APPIE OF
                                                                                                                                                                                                                   ARMITAGE, RICHARD
                                                                                                                                                                                                                                        FANSLOW, WILLIAM
SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                         ARVIA E.
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TOPOLOGY: linear
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CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: WASHINGTON
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                                                                                                         Indels
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APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSION, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                               Score 631.4; DB 3;
Pred. No. 3.1e-177;
                                                                                                         0; Mismatches
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99.8%; Pred. No. 3.1
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51 UNIVERSITY STREET
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                                                                                                         Matches 632; Conservative
             mat_peptide
93..857
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                                                                                           Similarity
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STATE: WASHING
COUNTRY: USA
; FEATURE:
; NAME/KEY: ;
; LOCATION:
US-09-088-913A-22
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                           COMPUTER: Apple Macintosn
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, version 5.1a
SOFTWARE: PAPLICATION DATA:
APPLICATION NUMBER: US/08/769,819
FILING DATE: 19-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           80.6%; Score 631.4; DB 4;
99.8%; Pred. No. 3.1e-177;
iive 0; Mismatches 1;
                                                                                                                                                                                                                       APPLICATION NUMBER: 08/249,189
FILING DATE: May 24, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING NAME: 07/805,723
                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802-E
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                               FILING DATE: 07-JUN-1995
APPLICATION NUMBER: 08/240 107
FILING DATE:
                                                                                                                                                                                                                                                                                                                                            WUMBER: 07/805,723
December 5, 1991
                                                                                                                                                                                                                                                                                                                                                                                                 07/783,707
                                                                                                                                                                                                                                                                                                                                                                                                                     FILLING DATE: October 25, 1991 ATTORNEY/AGENT INFORMATION:
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                 E: Floppy disk
Apple Macintosh
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SEQUENCE CHARACTERISTICS:
LENGTH: 878 base pairs
TYPE: nucleic acid
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Matches 632; Conservative
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93..857
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APPLICATION NUMBER:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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NAME/KEY:
LOCATION:
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; LOCATION:
US-08-769-819-22
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GGCCTCTGGCTGAAGCCCATTGGATCTGAAGAGTCTTAAGCCGCAAATACCCGCCAATTACCCGCCAAATACCCGCCTCTGGAGGAATCTTACTAAGCCGCCAAATACCGGCCTTTACTCCAGGCGCCAAATACCGGCCTTTACTCCAGGCGCGCAATTACCGGCCTTCGGCGGAGTGGTTTACTTTACTCTGGCGGAGTTTTACTCTTTACTCTTTACTTTACTTTGGCAATTACCGAGTTTCTTTACTTTACTTTGGCGAGTTTTACTTTACTTTACTTTACTTTGGCGGAGTTTTATACTTTGCAGTTTTACTTTGCAGCGCAGTGTTTACTTTGTTTACTTTGTTTACTTTGTTTATTCTTTGTTTATTCTTTGTTTTTGTTTTTTTT	631 CACAGTTCCTCCCAGCTTTGCGAGCAGTCTGTTCACTTGGGCGGAGTGTTTGAATTA [11]
	631 705 691 765 751 825

Search completed: May 30, 2002, 02:51:25 Job time: 11589 sec AR076926 Sequence AR078316 Sequence

AR085419 Sequence AR103375 Sequence

AR106246 Sequence AR169232 Sequence

AR171647 Sequence 127345 Sequence Sequence Sequence

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1 (bases 1 to 786)
Srinivasans. and Spriggs,M.K.
Methods of preparing soluble, oligomeric proteins
Patent: US 5716805-A 3 10-FEB-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA
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Sequence 3 from patent US 5716805.
187864
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168 c 168 q
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ORIGIN
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JOURNAL
FEATURES
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KEYWORDS
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187864
                                                                         May 30, 2002, 02:48:45 ; Search time 6499.83 Seconds (without alignments) 2530.567 Million cell updates/sec
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                                                                                                                                                                       1 ATGATCGAAACATACAACCA.....TTGGCTTACTCAAACTCTGA 786
                                                                                                                                                                                                                                                                            3595312
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                                                                              1797656 seqs, 10463268293 residues
                                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
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Maximum Match 100%
Listing first 45 summaries
                                                      nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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length: 2000000000
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AF013985 Rattus no AF344853 Macaca ne AF079105 Felis cat AF086711 Canis fam

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H.sapiens T H.sapiens m AF344841 Cercocebu

AF344859 Macaca mu

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Human CD40-

PAT 10-AUG-1998

Score 776.4; DB 6; Length 786; Pred. No. 1.7e-191;

98.8%; 99.2%;

Query Match Best Local Similarity

Description

Query Match Length DB

Score

Result

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Location/Qualifiers
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Armitage, R.J., Fanslow, W.C. and Spriggs, M.K.
Monoclonal antibodies to CD40 ligand, pharmaceutical comprising the same and hybridomas producing the same comprising the S961974-A 11 05-OCT-1999;
Location/Qualifiers
                                                                                                                                                      Length 840;
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Pred. No. 1.7e-191
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Armitage, R.J., Fanslow, W.C., Spriggs, M.K., Srinivasan, S., Gibson, M.G., Morris, A.E. and McGrew, J.T.
Recombinant soluble CD40 ligand polypeptide and pharmaceutical composition containing the same Patent: US 5962406-A 11 05-0CT-1999;
Location/Qualifiers
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Pred. No. 1.7e-191;
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                 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAAGTCCCCCGGTAGA
                                                               TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
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GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG
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Armitage, R.J., Fanslow, W.C. and Spriggs, M.K.
CD40 ligand polypeptide
patent: US 6087329-A 11 11-JUL-2000;
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185 c 175 g
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                                                                                    GIGACIGAICCAAGCCAAGTGAGCCAIGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780
                                                                                              AACAAAGAGGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
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                                                                                                                                                                                                                                                                                                                                                                                     Length 840;
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Pred. No. 1.7e-191
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Pred. No. 1.7
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AR169232
AR169232.1 GI:17907047
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1 (bases 1 to 840)
Armitage, R.J., Fanslow, W.C.,
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99.2%;
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Best Local Similarity 99.2
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Treatment of individuals exhibiting defective (Patent: US 6106832-A 7 22-AUG-2000;
Location/Qualifiers
1.0840
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Pred. No. 1.7e-191;
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Sequence 7 from patent US 6106832.
AR106246.1 GI:12820776
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Kehry, M. and Castle, B. B. Deil culture system comprising high density ligand
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Spriggs,M.K.; Armitage,R.J. and Fanslow,W.C.
Detection of mutations in a CD40 ligand gene
Patent: US 5565321-A 7 15-OCT-1996;
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                            Aruffo, A., Hollenbaugh, D. and Ledbetter, J. Soluble and its use in B cell stimulation Patent: US 5540926-A 1 30-JUL-1996;
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765	780		-FEB-1998		T	· · ·		Gaps 0;	60. 105	120 - 165	180 225	240 285	300	360 <u>.</u> 405 <u>.</u>	420 465	480	540
CCAGGTGCTTCGGTGTTTGTCAAT	TTCACGTCCTTTGCTTACTCAAA		linear PAT 04-		L. and Murphy, W.J.	acterized by		Length 840; Indels 0; Ga	ACTGCCCATCAGC	TATCACCCAAATGATTGGATCTGTG 	AAAGGAATCTTCAT 	AAAGATCCTTATCC	AGGATATAATGTTA 	STGATCAGAATCCT 	CTGTGTTACAGTGG	AAAATGGGAAACAG 	CCTTCTGTTCCAAT
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AGTATTTGAATTGC	GAGCCATGGCACTG(840 bp stent US 5674492.	0)) nslow,W.C. III,	preventing or treating c cells expressing CD40 is 5674492-A 1 07-OCT-19 Location/Qualifiers	1. 840 /organism="unknown" 185 c 175 g 2	8%; Score 776.4 2%; Pred: No. 1 0; Mismatch	AAACTTCTCCCCGAT 	TACTTACTGTTTTCC	atagaaggttggaca 	AAACGATACAGAGAT 	TTAAAAGCCAGTTTG TTAAAAGCCAGTTTC	AAGAAAACAGCTTTC 	TAAGTGAGGCCAGCA 	CCATGAGCAACAACI 	GACTCTATTATATCT
CAATCCATTCACTTGGGAGGAGTATTTGAATTGCA	GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGC 	CICIGA 786 CICIGA:831	167828 Sequence 1 from pat	167828 167828.1 GI:2829950 Unknown	Unknown. Unclassified. 1 (bases 1 to 840 Armitage, R.J., Far	Method of preventing or treating neoplastic cells expressing CD40 Patent: US 5674492-A 1 07-0CT-199 Location/Qualifiers	1840 /organis: 266 a 185	98.8%; Similarity 99.2%;); Conservative	1 ATGATCGAAACATACAACCAACTTCTCCCCGATCTGCGGCCACTGGAACTGCCCATCAGC	ATGAAAATTTTTATGTATTTACTTACTGTTTTCCT 	CTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAGGAATCTTCAT 	GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACAAGAGGAGAAGATCCTTATCC 	TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 	aacaaagaggagacgaagaaagaaaa'agctttgaaatgcaaaaaggtgatcagaatcct 	CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 	GCTGAAAAAGGATACTACACCATGAGCAACATGGTAACCCTGGAAAATGGGAAACAG 	CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
706 CAAT	721 GTGA 766 GTGA	781 CTCTGA 826 CTCTGA		ACCESSION IG VERSION IG KEYWORDS .	NISM NCE ORS		SOURCE BASE COUNT ORIGIN	Query Match Best Local Similarity Matches 780; Conser	1 ATG 46 ATG	61 ATG 106 ATG	121	181	241	301	361	421	481
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minidae; Homo:
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Hollenbaugh, D., Grosmaire, L.S., Kullas, C.D., Chalupny, N.J., Braesch-Andersen, S., Noelle, R.J., Stamenkovic, I., Ledbetter, J.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Submitted (27-AUG-1992) Diane L Hollenbaugh, Bristol-Myers Squibb PRI, Seattle, WA, 98121 3 (bases 1 to 879) Hollenbaugh, D.
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The human T cell antigen gp39, a member of the TNF gene family, a ligand for the CD40 receptor: expression of a soluble form of gp39 with B cell co-stimulatory activity
EMBO J. 11 (12), 4313-4321 (1992)
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The original submission [1] reported 934bp.
Location/Qualifiers
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                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1803)
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Submitted (28-UUL-1992) M. Spriggs, Immunex Res & Development
Corporation, 51 University Street, Seattle WA 98101, USA
2 (bases 1 to 1803)
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/cell_type="peripheral blood T-cell"
46 .831
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Pred: No. 1.6e-191;
0; Mismatches 6;
          mRNA
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/protein_id="CAA48077.1"
/db_xref="GI:38412"
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H.sapiens mRNA for CD40 ligand.
X67878 S50586
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                                                                                                     601 TICGAGAGAATCITACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCITGCGGGCAA
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GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

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Result
No.
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Maximum Match 100%
Listing first 45 summaries
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13736207 seqs, 6748477542 residues
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BF791160
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AZ166561 SP_0088_B
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BG715238 602676117
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ALIGNMENTS

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RESULT 'BF599437

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Contact: Smith TPL
COntact: Smith TPL
USDA, ARS, US Meat Animal Research Center
PO Box 166, Clay Center, NE 68933-0166, USA
Tel: 402 762 4366
Fax: 402 762 4390
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                                                                                         FORWARD: AGGAAACAGCTATGACCAT
BACKWARD: GTTTTCCCAGTCACGACG
Plate: 33 row: N column: 5
Seq primer: ATTTAGGTGACACTATAG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle Genome Res. 11 (4), 626-630 (2001)
                                                                                                                                                                                                                                          Email: smith@email.marc.usda.gov Single pass sequencing. Bases called and alt_trimmed with phred v0.980904.e. Vector identified by cross_match with the -minscor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
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/organism="Bos taurus'
/db_xref="taxon:9913"
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                                                           Sequence evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle Genome Res. 11 (4), 626-630 (2001)
                                                                                                                Pertea,G., Holt,I.,
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USDA, ARS, US Meat Animal PO Box 166, Clay Center, N
                             Contact: :Smith TPL
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                                                          Res. 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="MARC/
/tissue_type="poc
                                                                                                                                                                                        Bovinae;
                                                                                                                                                                                                       Eutheria;
                                                                                                                                                                                                     Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
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                                                                                                                                                              Roberts, A.J., Stone, R.T.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     233 GAAGATTTTGTGTTCATGAAAACGATACAGAGATGCAATAAAGGAGAGAGGGGTCCTTATCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTTTTTGCTGTGTATCTTCACAGAAGATTGGACAAGATAGAAGACGAAAGGAATCTTCAT 232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTTTTTGCTGTGTATACATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GAAGATTTTGTATTCATGAAAACGATACAGAGAGATGCAACACAGGAGAAAGATCCTTATCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Single pass sequencing. Bases call v0.980904.e. Vector identified by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and trimmed with phred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Plate: 34 row: F column: 23
Seq primer: ATTTAGGTGACACTATAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BACKWARD: GTTTTCCCAGTCACGACG
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           Tirunagaru, V.G., Sofer, L., Cui, J. and Burnside, J. an expressed sequence tag database of T-cell-enriched chicken splenocytes: sequence analysis of 5251 clones
                                                                            Archosauria; Aves; Neognathae; Phasianinae; Gallus.
                                                                                                                              Gallus gallus
                                                                                                                                                                              AI982044
AI982044.1
                                                                                                                                                                                                           pat.pk0072.c9.f chicken of clone pat.pk0072.c9.f 5'
                                                                                                                                                                                                                                          AI982044
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                                                                                                                Eukaryota; Metazoa;
                                                                                                                                                 chicken
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /tissue_type="pooled"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /db_xref="taxon:9913"
/clone_lib="MARC 1BOV"
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Library made from pooled tissue from
Library made from pooled tissue from
fat, hypothalamus, and pituitary."
fat, 77 c 82 g 105 t
66 (2), 144-151 (2000)
                                                                                                                                                                                GI:5885072
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                                                                                                               Chordata;
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                                                                                                                                                                                                             638 bp mRNA activated T cell similar to CD40
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                                                                                                                                                                                                              linear EST 07-MAY-2001 cDNA Gallus gallus cDNA ligand, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                         SP_0088_Bl_C02_T7A Strongylocentrotus purpuratus, sperm genomic BAC library Strongylocentrotus pur clone Plate=88 Col=3 Row=F, DNA sequence.
AZ166561
                                                                                                                                                                                                                                   Strongylocentrotus purpuratus
Eukaryota; Metazoa; Echinodermata; Eleutherozoa;
Echinoidea; Euechinoidea; Echinocea; Echinoida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Contact: Joan Burnside
Molecular Endocrinology
                                                                                                                                          Cameron,R.A., Mahairas,G., Rast,J.P., Martinez,P., Biondi,T.R., Swartzell,S., Wallace,J.C., Poustka,A.J., Livingston,B.T., Wray,G.A., Ettensohn,C.A., Lehrach,H., Britten,R.J, Davidson,E.H. a
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                                                                                                                                                                                                                                                                                        Strongylocentrotus purpuratus.
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                  Division of Biology 156-29
                                                                    Proc. Natl. Acad.
                                                                                                          A sea urchin genome project: Sequence scan,
                                                                                                                                                                                                                  Strongylocentrotidae; Strongylocentrotus
                                                                                                                                                                                                                                                                                                                              AZ166561.1 GI:8336929
                                  Contact: Cameron, RA, Davidson, EH,
                                                      20402566
                                                                                          additional
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primer: T7.
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/clone="pat.pk0072.c9.f"
/clone_lib="chicken activated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /cell_type="Con A-activated
/lab_host="E.coli TOP10 F'"
/note="Vector: pcDNA3"
.171 c 151 g 180 t
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of Technology
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Pred. No. 1.4e-05;
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ACCESSION
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                                                                                                                                                                                                                                                                                                                                                                                             CNS005TE 997 bp DNA linear GSS 0 Drosophila melanogaster genome survey sequence TET3 end o BACK12K22 of RPCI-98 library from Drosophila melanogaster
                  collaboration with the Berkeley Drosophila Genome Project (BDGP). The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further information please see http://www.fruitfly.org The BDGP Drosophila melanogaster BAC library was prepared by Kazutoyo Osoegawa and

    Web: www.genoscope.cns.fr)
    Determination of this BAC-end sequence was carried out as

                                                                                                                                             Submitted (02-JUN-1999) Genoscope - Centre National de Sequencage BP 191 91006 EVRY cedex - FRANCE (E-mail : segref@genoscope.cns.fr
                                                                                                                                                                                                                                                         Eukaryota;
Pterygota;
                                                                                                                                                                                                                                                                                                                                                               fly), genomic survey sequence. ALO60767
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Location/Qualifiers
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Class: BAC ends
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: acameron@caltech.edu
Plate: 88 row: F column:
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erygota; Neoplera; Endopterygota; Diptera; Brachycera;
scomorpha; Ephydroidea; Drosophilidae; Drosophila.
(bases 1 to 997)
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(626) 793-3047
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/db_xref="taxon:7668"
/clone="Plate=88 Col=3 Row=F"
/clone=!be"Strongylocentrotus purpuratus,
urchin, sperm genomic BAC library"
/note="Organ: sperm; Vector: BACe3.6; BAC
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Submitted (23-JUL-1999) Genoscope
BP 191 91006 EVRY cedex - FRANCE (
- Web : www.genoscope.cns.fr)
Determination of this BAC-end sequ
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1 (bases I to 1043)
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/clone_lib="RPCI-98"
/clone="BACR12K22"
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Pred. No..0
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            sequence was carried out as
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                                                                     (E-mail -:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ТАМАМААААНЫМАМАМАМАЛААНЫТАТТТИМИНАМАТАААМАМАМИМИМАААААНАТАТА 617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            113;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      http://www.edgp.ebi.ac.uk . This Drosophila melanogaster BAC library (Dros BAC) was made by Alain Billaud at CEPH (Centre d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     pBeloBAC11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           collaboration with the European Drosophila Genome Project http://www.edgp.ebi.ac.uk - This Drosophila melanogaster
                                                                                                                                                                                                                                                                                                                                                                        MR1-RT0025-171100-003-c06 RT0025
BF820152
                                                                                                                                           Goldman,G.H., Carvalho,A.F., Matsukuma,A., Baia,G.S., Simpson,D.H. Brunstein,A., deoliveira,P.S., Bucher,P., Jongeneel,C.V., O'Hare,M.J., Soares,F., Brentani,R.R., Reis,L.F., de Souza,S.J. and
                                                                                                                                                                                                   1 (bases 1 to 396)
Dias Neto, E., Garcia Correa, R., Verjovski-Almeida, S., Briones, M.R.
Nagai, M.A., da Silva, W. Jr., Zago, M.A., Bordin, S., Costa, F.F.,
                                                                                                                                                                                                                                                                                                                                                   BF820152.1
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Contact: Simpson A.J.G.
Laboratory of Cancer Ge
Ludwig Institute for Ca
                                                                                                          Shotgun sequencing of
                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                         sequence tags
                                                                                                                          Simpson, A.J.
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/note="end : T7"
/note="end : T7"
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/plasmid="pBeloBAC11"
/db_xref="taxon:7227".
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35.1%;
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                     Genetics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             266 CAGAAAAATGTAGAAGTGAAAAACGAAAAAGAACAAACAGAAAAACCCTGGAAATGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              238 TCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  86 TTCATACATTACAGTGGTTGGAATAAAAATTGGGATGAATGGGTTCCGGAGAGCAGAGTA 145
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
nes 118; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CAGTATGCAGAGGGGAAGATGAGAGGGGGCTGCCCCAGGAAAGAAGACATCTGGTCTGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CCTCAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAG 417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TTAAACAAAGAGAGACGAAGAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAT 357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TGGGCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGG 473
                                                                                                      ,G., Cheng,Z. and Han,Z.
Homo sapiens cDNA DCB clones
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                           AV714252 DCB Homo sapiens cDNA
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                                                                                                                                                                                                                                                                                                                                                                                                           AV714252
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                Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park,
201203, P. R. China
                                                                                      Contact: Zeguang Han
                                                                                                                                                                         Xu,X., Gu,J., Liu,F., Qu,J., Zhao,M., Li,Y., Huang,Q., Zhou,J., Song,H., Gu,Y., Yang,Y., Gao,G., Xiao,H., Li,N., Qian,B., Gao,X., Cheng,Z., Xu,S., Gu,W., Tu,Y., Jia,J., Fu,G., Ren,S., Zhong,M., Lu
                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Primates; 1 (bases 1 to 486)
                                                                                                                                                                                                                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (http://www.ludwig.org.br/scripts/gethtml2.pl?t1=MR1&t2=MR1-RT0025-
771100-003-c06&t3=2000-11-17&t4=1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: asimpson@ludwig.org.br
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  primer: puc 18 forward
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86-21-50801919(ex.45)
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Site_2: SmaI; A mini-library was made by cloning products
derived from ORESTES PCR (U.S. Letters Patent application
No. 196,716 - Ludwig Institute for Cancer Research)
profiles into the pUC 18 vector. Reverse transcription or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
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50.0%;
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Pred. No. 1.5;
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clone DCBADF07 5',
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                                                                                                                                                                                                                                                                                                                      Gu,Y., Peng,Y., Souy,.... Qu,J., Gau
Li,N., Qian,B., Liu,F., Qu,J., Gau
Li,N., Gi,W., Tu;Y., Jia,J., Fu,G.,
,S., Gu,W., Tu;Y., Jia,J., Fu,G.,
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AV724813
AV724813
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Email: hanzg@chgc.sh.cn
This clone is available at CHGC in
                                                                                                                                                                                                                                   Contact: Zeguang Han
Chinese National Human Genome Center at Shanghai
Chinese National Road, Zhangjiang Hi-Tech Park, Pudong,
                                                                                                                                                                                                                                                                                                    Homo sapiens cDNA HTB clones 
Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
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                                                                                                                                                                                                           201203, P. R. China
Tel: 86-21-50801919(ex.45)
                                                                                                                                                                                              Fax: 86-21-50801922
                                                                                                                                                                                                                                                                                                                                                                                                               (bases 1 to 578)
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                                                                                                                                                             clone
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                                                                                                                                                       hanzg@chgc.sh.cn
lone is available at CHGC
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87 c 124 g 93 t
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/cell_type="dendritic cells"
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              /tissue_type="Hypothalamus"
/dev_stage="Adult"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /dev_stage="mature"
/lab_host="BM25.8"
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                                                 /clone_lib="HTB"
                                                                  /clone="HTBAKC04"
                                                                                     /organism="Homo sapiens"
/db_xref="taxon:9606"
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Pred. No. 1.5;
0; Mismatches 118;
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3., Ren,S.,
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HTBAKC04 5',
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S., Zhong,M., Lu,
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mRNA sequence.
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G., Hu,R.,
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                                                                                                                                                                   source
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (2000)
Other_ESTs: 1e02f03.x1
Contact: Douglas Melton,
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Hillier,L., Marra,M., Pape,D., Wylle,T., Martin,J., Blistain,A.,
Schmitt,A., Theising,B., Ritter,E., Ronko,I., Bennett,J., Carden
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                                                                                                                                                                                    High quality sequence stop: 47.
Location/Qualifiers
                                                                                                                                                                                                                                                                  obtaining a clone please contact: Dr. Hiroshi Inoue
                                                                                                                                                                                                                                                                                       Library was constructed by Dr. Hiroshi Washington University Genome Sequencing
                                                                                                                                                                                                                                                                                                                                                                                                           Harvard University, Howard Hughes Medical Dept of Molecular and Cellular Biology, 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                              indocrine Pancreas Consortium
                                                                                                                                                                                                                                                   (hinoue@im.wustl.edu)
                                                                                                                                                                                                                                                                                                                               mail: dmelton@biohp.harvard.edu

    Gibbons, M., McCann, R., Cole, R., Tsagareishvili, R., Williams, T. Jackson, Y. and Bowers, Y.

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617-495-8557
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Not1; Site_2: XhoI; cDNA made by oligo-dT priming. Size-selected on agarose gel. Average insert size
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                         /note="Organ:
NotI; Site_2:
                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone_lib="HR85 islet"
                                                               /tissue_type="Purified pancreatic islet"
/lab_host="DH10B"
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BF791160
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602251255F1 NIH_MGC_81 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs-r@mail.nih.gov
Tissue Procurement: CLONETECH Laboratories, Inc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EST
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Contact: Robert Strausberg, PhiD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Ve
Mammalia; Eutheria; Primates; Catarrhin1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                   tound
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CDNA Library Preparation: CLONETECH Laboratories, I CDNA Library Arrayed by: The I.M.A.G.E. Consortium DNA Sequencing by: Incyte Genomics, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                               nttp://image.lini.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                Clone distribution: MGC clone distribution information can be cound through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (bases 1 to 586)
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                                                                                                                                                                                                                                                                                                                                     quality sequence stop: 582
Location/Qualifiers
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Amplified once. Contact information: Hiroshi Inoue, I
Metabolism Div. (Alan Permutt Lab), Washington Unive.
School of Medicine, Box 8127, 660 South Euclid Ave.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             314-362-1916,
101 c
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5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3' (where B = C, or G and N = A, C, G, or T). Average insert size 1.55 kb (range 1.0-4.0 kb). 15/15 colonies contained
                                                                            (ggccattatggcc); 5' and 3' adapt as follows: 5' adaptor sequence: and 3' adaptor sequence:
                                                                                                /lab_host="DH10B (TI phage-resistant)"
/note="Organ: muscle (skeletal); Vector: pDNR-LIB
/note="Organ: muscle (skeletal); Vector: pDNR-LIB
(Clontech); Site_1: Sfil (ggccgcctcggcc); Site_2: Sfil
(ggccattatggcc); 5' and 3' adaptors were used in cloning
as follows: 5' adaptor sequence: 5'-CACGGCCATTATGGCC-3'
                                                                                                                                                                                                                                                                  /organism="Homo sapiens
/db_xref="taxon:9606"
                                                                                                                                                                                                                      /clone_lib="NIH_MGC_81"
                                                                                                                                                                                                                                            /clone="IMAGE:4338636"
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Pred. No. 1.5;
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inserts by PCR. This library was enriched for full-length clones and was constructed by Clontech Laboratories (Palo

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REFERENCE
AUTHORS
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ORGANISM
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118; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata;
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                                                                                                                                                                                  /note="Organ: Fetal brain; Vector: pCMVSPORT 6; 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the pcMVSPORT 6 vector. Library was constructed by Life
                                                                                  http://fulllength.invitrogen.com"
43 c 120 g 152 t 7
                                                                                                                                   Technologies. Contact: Feng Liang Life Technologies; division of Invitrogen 9800 Medical Center Drive Rockv:
, Maryland 20850, USA Fax: (1) 301 610 8371 Email:
                                                                                                                                                                                                                                                                       /clone_11b="LTI_FL013_FBrn1".
/dev_stage="pooled tissue from post conception fetuses (20 week, 24 week and 26 week)"
/lab_host="DH10B"
                                                                                                                      fliang@lifetech.com URL :
                                                                                                                                                                                                                                                                                                                                          /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="CSODF004YD24"
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Pred. No. 1.5;
O; Mismatches
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     Score 47.2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGCCAGCAGTAAAACAACATCTGTGTTACAGTGGGCTGAAAAAGGATACTACACCATGAG 446
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAATRTAAAAAAAAAAAAAAGTAAATTWAWTTWAWRRAGGATAAAAAATAAAAAAAATGRA 363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AGATGCAACACAGGAGAAAGATCCTTATCCTTACTGAACTGTGAGGAGATTAAAAGCCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         168;
                                                                                                                                                                                                                                                              Email: cgapbs-r@mail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
cDNA Library Preparation: Michael J. Brownstein (NHGRI), Sh
Toshiyuki and Piero Carninci (RIKEN)
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incytte Genomics, Inc.
Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                                                                                     NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian
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                                                                                                                                                                                                         http://image.llnl.gov
Plate: LLAM10686 row: b column:
                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                      Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BG715238.1
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                                                                                                                                                                                     High quality sequence stop: 830
                                                                                                                                                                                                                                              found through the I.M.A.G.E. Consortium/LLNL
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/lab_host="DH10B"
/note="Organ: brain; Vector: pBluescriptR (modified
pBluescript KS+); Site_1: BamHI; Site_2: SalI-XhoI
                                                                     /clone="IMAGE:4798372"
/clone_lib="NIH_MGC_96"
                                                       /tissue_type="hypothalamus"
                                                                                                            /organism="Homo sapiens"
/db_xref="taxon:9606"
                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Metazoa;
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Primates;
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                                                                        _96"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       238 TCCTTACTGAACTGTGAGGAGATTAAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 298 TTAAACAAAGAGGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAT 357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TTCATACATTACAGTGGTTGGAATAAAAATCGGGATGAATGGGTTCCGGAGAGCAGAGTA 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CAGTATGCAGAGGGGAAGATGAGAGGGGGCTGCCCCAGGAAAGAAGACATCTGGTCTGCAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    118;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Tissue Procurement: Miklos Palkovits, M.D., cDNA Library Preparation: Michael J. Browns Toshiyuki and Piero Carninci (RIKEN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mRNA
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BI601432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Unpublished (1999)
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                                                                                                                                                                                                                                                                                                                   http://image.llnl.gov
Plate: LLAM11761 row:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         National Institutes of Health,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NIH-MGC http:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mammalia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                               found
                                                                                                                                                                                                                                                                                                                                                                                  DNA Sequencing by: Incyte Genomics, Inc. Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                            cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
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                                                                                                                                                                                                                                                                                           quality sequence stop:
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                                                                                                                                                                                                                                                                                                                                                          through the I.M.A.G.E. Consortium/LLNL at:
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size-selected for average
normalized to ROT 5. This i
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."
pBluescript KS+); Site_1: BamHI; Site_2: SITe_XII-XhoI (gtcc); Oligo-dr primed using primer 5'-TTTTTTTTTTTTTTTTVN-3' size-selected for average insert size 2.3 kb and normalized to ROT 5. This is a primary library enriched
                                                                                                                                                        /clone="IMAGE:5300787"
/clone_lib="NIH_MGC_96"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for full-length clones and constructed using the
                                                                                           /lab_host="DH10B"
/note="0---
                                                                                                                                                                                                  /organism="Homo sapiens"
/db_xref="taxon:9606"
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                                                                                      note="Organ: brain; Vector: pBluescriptR (modified
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         M.D., Ph.D.
Brownstein
                                                                 Site_2: SalI-XhoI
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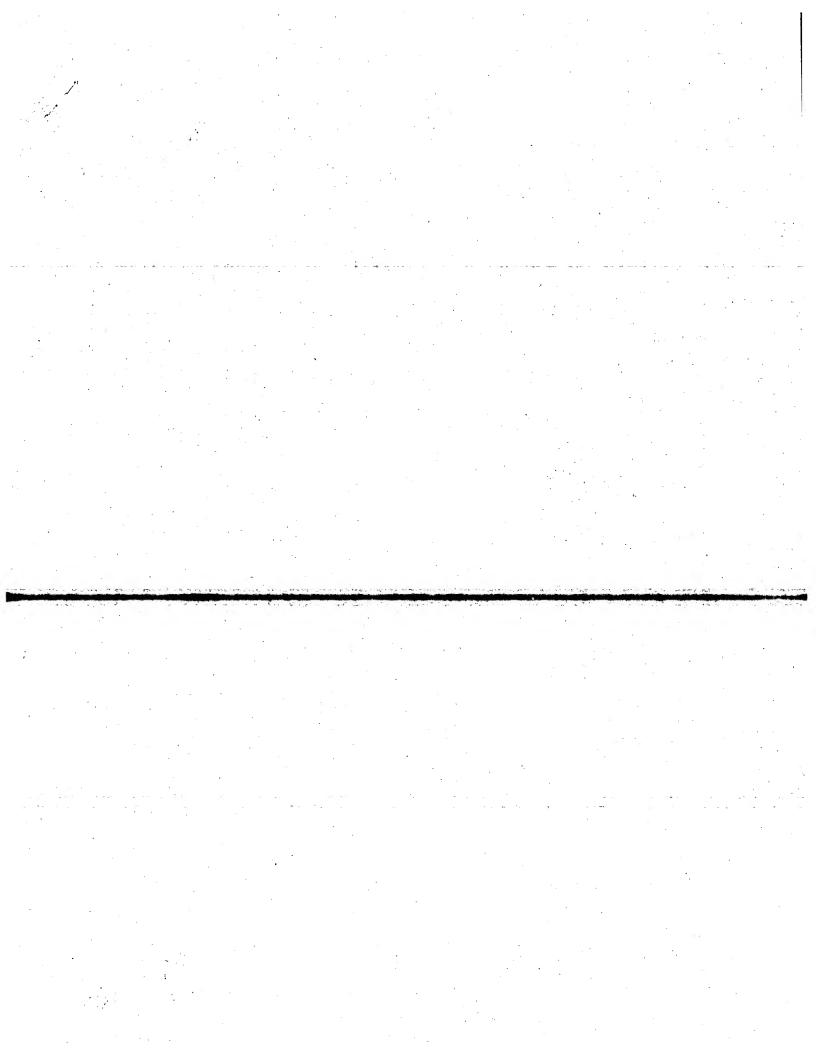
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AUTHORS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CAGTATGCAGAGGGGAAGATGAGAGGGGCTGCCCCAGGAAAGAAGACATCTGGTCTGCAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BF573761 861 bp mRNA linear EST 12-DE 602077072F2 NIH_MGC_62 Homo sapiens cDNA clone IMAGE:4251880
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BF573761
                                                                                                                                                                                                                                                                                                                                                             High quality sequence stop:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Unpublished (1999)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BF573761.1 GI:11647473
                                                                                                                                                                                                                                                                                                                                                                                  Plate: LLCM1075
                                                                                                                                                                                                                                                                                                                                                                                                     http://image.linl
                                                                                                                                                                                                                                                                                                                                                                                                                       found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Tissue Procurement: ATCC/DCTD/DTP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Email: cgapbs-r@mail.nih.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        National Institutes of Health, Mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NIH-MGC http://mgc.nci.nih.gov/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cDNA Library Preparation: CLONETECH Laboratories, Inc. cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                      DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can
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Sfil (ggccgcctcggcc); Site_2: Sfil (ggccattatggcc); Double-stranded cDNA was prepared from cell line RNA. 5 and 3' adaptors were used in cloning as follows: 5' adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor sequence: 5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cap-trapper method (Carninci, in preparation). Li constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Lib 145 c 243 g 178 t
                                                                                                                                                                              /tissue_type="melanotic melanoma, high MDR"
/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: skin; Vector: pDNR-LIB (Clontech); Site_1:
                                                                                                                                                                                                                                           /clone="IMAGE:4251880"
/clone_lib="NIH_MGC_62"
                                                                                                                                                                                                                                                                                  /db_xref="taxon:9606
                                                                                                                                                                                                                                                                                                   organism="Homo sapiens"
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Search completed: May 30, 2002, 01:00:07 Job time: 22217 sec



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Title:
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Sequence:
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Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Gapop 10.0 , Gapext 1.0
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ld-geneseq/geneseqn-embl/NA1995 DAT:
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Human CD40 ligand
cDNA of CD40L, a n
Human CD40 ligand
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WPI; 1998-348521/30.		Cantwell M. Kipps T.J. Sharma S:	(REGC) UNIV CALIFORNIA.			01-DEC-1997; 97US-0982272.		08-DEC-1997; 97WO-US22740.		18-JUN-1998.		WO9826061-A2.		Chimeric - Homo sapiens.	Chimeric - Mus sp.		autoimmune disorder; rheumatoid arthritis; vaccine; chimera; ss.	accessory molecule ligand; AML; gene therapy; treatment; neoplasia;	CD40 ligand; alteration; immunoreactivity; human cell;		Exemplary CD40 ligand gene used in the course of the invention.		23-SEP-1998 (first entry)		AAV39000;	ANYSSUOD SCAHAALA, DNA, 100 BE.	20000 1152222 705	

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cytoplasmic domain (Domain I). The sequence is used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells. Vectors containing the AML genes can be used in gene therapy for treating nephasia or autoimmune disorders such as rheumatoid arthritis. They can also be used for vaccination to produce immunity against a virus cell, bacteria, protein, fungus or neoplasia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents an exemplary CD40 ligand gene, comprising nucleotides encoding the extracellular domains (Domains I and IV) of human CD40 ligand gene (AAV38998) are operatively linked nucleotides encoding the murine CD40 ligand transmembrane domain (Do 1) (AAV38997) which is operatively linked to the human CD40 ligand grant l
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure;
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                                                                                                                                       A DNA fragment encoding the extracellular (soluble) region of human CD400-L was ligated to a synthetic oligonucleotide sequence coding for a leader peptide, a 3 amino acid leucine zipper sequence (AAR53968) and the Flag (RTM) linker sequence. Cells expressing the fusion construct are grown to accumulate oligomeric, soluble CD40-L fusion construct are grown to accumulate oligomeric.
                                                                                                                                                                                                                                            Prepn. of soluble oligomeric mammalian proteins cells to express a fusion protein comprising a l domain and a heterologous mammalian protein
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                                                                                                                                                                                                                                                                                                    caatccattcacttgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaat
                                                                                                                                                                                                                                                                                                                                                     ttcgagagaatcttactcagagctgcaaatacccacagttccgccaaaccttgcgggcaa
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                                                                                                                                          standard;
                                   molecule
                       disorder;
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                                                                   gene used in
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                                             alteration;
                                                                                           (first
                                 ligand;
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                                                                                           entry)
                       rheumatoid
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                                                                    the
                                             immunoreactivity;
                                   AML;
                                                                   course of the
                      gene therapy; arthritis; va
                       vaccine; ss
                                             human cell;
                                                                    invention
                                treatment;
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Best Local
                                                                                                                                                                The present sequence represents the CD40 ligand gene. The sequence is used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells. Vectors containing the AML genes can be used in gene therapy for treating neoplasia or autoimmune disorders such as rheumatoi arthritis. They can also be used for vaccination to produce immunity against a virus cell, bacteria, protein, fungus or neoplasia.
                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                         Vectors containing accessory molecule ligand genes. used for altering immunoreactivity of cells, particularly for treatment neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                             01 DEC-1997;
09-DEC-1996;
                                                                                                                                                                                                                                                                                                  Disclosure;
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  61
                                       ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGCCCACTGGACTGCCCATCAGC
  ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG
                           atgatcgaaacatacaaccaaacttctccccgatctgcggccactggactgcccatcagc
                                                                               780;
                                                                                          Similarity
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                                                                              Conservative
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96US-0032145
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Pred. No. 6.2e
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tes 6;
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Š 밁 Qy 밁 Ş 밁 Ş DЬ Ş В Q 밁 δÃ 밁 421 481 421 361 361 301 301 241 241 181 181 121 121 61 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT aacaaagaggagacgaagaaagaaaacagctttgaaatgcaaaaaggtgatcagaatcct AACAAAGAGGAGGACGAAGAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA gaagattttgtattcatgaaaacgatacagagatgcaacacaggagaaagatccttatcc GAAGATTTTGTATTCATGAAAAACGATACAGAGATGCAACAGAGAGAAAGATCCTTATCC ctttttgctgtgtatcttcatagaaggttggacaagatagaagatgaaaggaatcttcat GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG ttactgaactgtgaggagattaaaagccagtttgaaggctttgtgaaggatataatgtta 300 180 480 480 420 420 240 180 120 540 360 360 300 240 0

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25-OCT-1991;
05-DEC-1991;
23-OCT-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Leucine zipper; fusion protein production; soluble oligomeric protein; heterologous mammalian type II transmembrane protein; activated T cell; heterologous mammalian type I transmembrane protein; antibody production; CD40-L; B-cell proliferation; CD27-L; lymphocyte antigen; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CD40 ligand coding sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAV12852;
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                                                                                                                                         WPI; 1998-144799/13.
P-PSDB; AAW41178.
                                                                                                                                                                              Spriggs
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This sequence is the coding sequence for the human CD40 ligand (CD40-L). The encoded protein can be used in a fusion protein produced using the method of the invention. The method is for preparing soluble oligomeric protein by culturing a host cell transfected with a vector for the fusion protein. The soluble oligomeric proteins comprise a leucine zipper
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                                                                                                               soluble oligomeric fusion proteins
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                                                                                                    extracellular region of
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91US-0783707.
91US-0805723.
92US-0969703.
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                                                                            21pp; English.
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protein, where the leucine zipper is a peptide comprising at least part of AAW41171 or AAW41172, optionally with conservative amino acid substitutions, provided that the peptide trimerises in solution. A substitutions, provided that the peptide trimerises in solution activated soluble fusion protein comprising the leucine zipper of AAW41171 linked to the extracellular region of CD40-L (a type II transmembrane protein that is found on activated T cells and acts as a ligand for the B-cell antigen CD40) stimulates B-cell proliferation and antibody production in a similar manner to membrane-bound CD40-L. A soluble fusion protein comprising the leucine zipper of AAW41172 linked to the extracellular region of CD27-L (a type II transmembrane protein that binds to the leucine cD27) inhibits binding of CD27c (a fusion protein lymphocyte antigen CD27) inhibits binding of CD27c (a fusion protein lymphocyte antigen CD27) inhibits binding of CD27c (a fusion protein lymphocyte antigen CD27) inhibits binding of CD27c and a human IgG1 Fc region) extracellular region of a heterologous mammalian type I trans protein, where the leucine zipper is a peptide comprising at fused mammalian type II Ġ EBV-transformed B cells expressing to the N terminus of the extracellular region of a hete lian type II transmembrane protein or to the C terminus CD27-L. I transmembrane a heterologous rminus of the reast part in

Sequence 786 BP; 250 A; 168 C; 168 G; 200 T; 0 other;

Length

786;

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g QY 2 Вþ Qy Д δÃ B Вþ QΥ В δÃ 밁 QΥ QΥ 밁 Ş δÃ В ρ Qy 멍 Ω В Best Local Sir Matches 780; Query Match Best Local S 241 181 181 241 121 121 541 541 481 481 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT 421 421 361 361 301 301 AACAAAGAGGAGAGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 61 61 661 601 601 661 1 atgatogaaacatacaaccaaacttctccccgatctgcggccactggactgcccatcagc 1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCCACTGGACTGCCCATCAGC atgaaaatttttatgtatttacttactgtttttcttatcacccagatgattgggtcagca 120 ATGAAAATTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG caaattgcggcacatgtcataagtgaggccagcagtaaaacaacatctgtgttacagtgg TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300 gaagattttgtattcatgaaaacgatacagagatgcaacacaggagaaagatccttatcc GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC ctttttgctgtgtatcttcatagaaggttggacaagatagaagatgaaaggaatcttcat CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG aacaaagaggagacgaagaaagaaaacagctttgaaatgcaaaaaggtgatcagaatcct ttactgaactgtgaggagattaaaagccagtttgaaggctttgtgaaggatataatgtta ctgaccgttaaaagacaaggactctattatatctatgcccaagtcaccttctgttccaat gctgaaaaaggatactacaccatgagcaacaacttggtaaccctggaaaatgggaaacag GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG TTCGAGAGAGATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA cgggaagcttcgagtcaagctccatttatagccagcctctgcctaaagtcccccggtaga CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT ttcgagagaatcttactcagagctgcaaatacccacagttccgccaaaccttgcgggcaa caatccattcacttgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaat Similarity Conservative 98.8%; 0; Score 776.4; DB 19; Pred. No. 6.2e-214; Mismatches 6; Gaps 240 180 180 120 60 60 240 480 420 420 360 360 300 600 540 540 480 780 660 600 720 660 720

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RESULT
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Best Local Similarity
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                                                                                                                   binding with CD40 an dinduces B cell proliferation and induction of antibody secretion, including IgE. Membrane bound CD40-L acts as a CD40 agonist and soluble CD40-L acts as a CD40 antigonist. CD40-L acts as a CD40 antigonist. CD40-Can be used in a binding assay to detect cells expressing CD40.
                                                                                                                                                       This sequence encodes a human CD40-L polypeptide which binds to CD40. CD40-L is a type II membrane polypeptide which has an extracellular region at its C-terminus, a transmembrane region and an intracellular region at its N-terminus. A soluble form of CD40-L lacks the transmembrane domain. CD40-L activity is mediated by
                                                                                                                                                                                                                            New cytokine CD40-L as CD40 agonist and antagonist - is used for treating allergies, lupus, rheumatoid arthritis, graft-versus-host disease and insulin-dependent diabetes mellitus
                                                                                                    Sequence 840
                                                                                                                                                                                                             Claim 1; Fig
                                                                                                                                                                                                                                                                                             Armitage RJ,
                                                                                                                                                                                                                                                                                                               (IMMV ) IMMUNEX CORP.
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05-DEC-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     proliferation; inducantagonist; binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; CD40-L;
transmembrane;
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DB; AAR36701.
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ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG
                   atgatcgaaacatacaaaccaaacttctccccgatctgcgccactggactgcccatcagc
                                                       Conservative
                                                                                                    BP;
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91US-0805723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               induction; antibody; secretion;
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                                                               98.8%;
99.2%;
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                                                     Score 776.4; DB 14
Pred. No. 6.4e-214;
0; Mismatches 6;
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                                     CDS
                                                                               Probe; primer; mutation; CD40
W09417196-A
                                                               Synthetic
                                                                                         Probe;
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                                                                                                                              23-MAR-1995
                                                                                                                                                AAQ67123;
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CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTTGTCAAT
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                                           GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
                                                                                  caatccattcacttgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaat
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PCR; amplify; ligand gene; Location/Qualifiers 46..831 /*tag= a /*tag= a /product= CD40 ligand DNA; 840 igm; ss. ВP

chain

reaction;

detection;

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22-JAN-1993;
20-JAN-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence represents the CD40 ligand gene. Mutations within this gene were identified by the method of the invention. The method comprises isolating DNA from an individual and selectively amplifing the isolated DNA derived from the CD40 ligand gene. The amplification product is then analysed to determine if there is a mutation present and determining if a protein expressed from the ligand gene will bind CD40. The detection of mutations in the CD40 ligand gene allows subsequent treatment of a syndrome resulting in elevated lavels of serum IgM and diminished levels of other Ig isotypes, due to mutation in the CD40 ligand gene. ie. X-linked.
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DB; AAR57469.
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                                                                                                            CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
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                                                                                                                                                              aacaaagaggagacgaagaaagaaacagctttgaaatgcaaaaaggtgatcagaatcct
   CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                               gctgaaaaaggatactacaccatgagcaacaacttggtaaccctggaaaatgggaaacag
                                                GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG
                                                                                               caaattgcggcacatgtcataagtgaggccagcagtaaaacaacatctgtgttacagtgg
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The nucleotide sequence given in AAT05763 encodes a human high-density, membrane-bound (hdmb) CD40 ligand (AAR85486) that induces long-term proliferation of B-cells in culture. These proliferation B-cells proliferation of the differentiate into antibody-prodg. cells. The can be induced to differentiate into antibody-prodg. cells. The nucleotide sequence is incorporated into a baculovirus vector that

nucleotide sequence is incorporated in is used to transfect Sf9 insect cells

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RESULT
AAT05763
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                        High density membrane bou
proliferation of B cells
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P-PSDB; AAR85486.
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Disclosure; Fig 1; 74pp; English
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                                            bound CD40 ligand - for stimulat
lls in vitro or in vivo, partic.
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Query Match Best Local S Matches 780

Similarity

98.8%; 99.2%;

Conservative

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Mismatches

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ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCCACTGGACTGCCCATCAGC

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                   (first
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novel cytokine
                                                           cDNA;
                   entry)
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 ligand for CD40
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The present sequence represents the cDNA sequence of a novel cytokine ligand for CD40 called CD40L CD40L is a type II membrane polypeptide, which is expressed by activated CD4+ T cells, and causes B cell proliferation and induction of antibody secretion. The protein can be used to produce monoclonal antibodies, which in turn bind to CD40-expressing cells. This inhibits binding of soluble CD40 to its ligand CD40L. The monoclonal antibody against CD40L is used to inhibit proliferation of neoplastic cells, and is particularly useful in treating B-cell lymphoma (e.g. where induced after transplants or in other cases of immune deficiency such as AIDS), and also melanoma or carcinoma. Since the monoclonal antibodies inhibit neoplastic cells directly, they may not need to be coupled to a toxin or radioisotope, avoiding toxic effects on normal B cells.
                                                                                                                                                                                                                                     Treating or preventing neoplastic disease associated with CD40-expressing cells - particularly B-cell lymphoma, by administration of CD40-binding protein, preferably antibosoluble CD40-ligand
                                                                                                                                                                                                                                                                                                                    WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CDS
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                                                                                                                                                                                                             Claim 3;
                                                                                                                                                                                                                                                                                                                                            Armitage
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               melanoma; carcinoma;
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Sequence
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DB; AAW34669.
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 840
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  B₽;
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HEALTH & HUMAN SERVICES
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46..831
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B-cell 1
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a; immune deficiency;
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98.8%; 1; DB 18; 5.4e-214; Length 840;

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                                   GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGAGAGAAGAATCCTTATCC
                                                                        CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT
                           gaagattttgtattcatgaaaacgatacagagatgcaacacaggagaaagatccttatcc
                                                             ctttttgctgtgtatcttcatagaaggttggacaagatagaagatgaaaggaatcttcat
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0; Mismatches
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                                                    28-APR-1995;
28-APR-1994;
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           Castle B,
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                              (BOEH ) BOEHRINGER INGELHEIM
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                                                                                                                                                                                                                                                                      CD40 ligand encoding
                                                                                                                                                                                                                                     CD40 ligand; TNF receptor family; activated T cell; membrane glycoprotein; cell proliferation; differen
                                                                                                                                                                                                                                                                                                                                                                                         831
           Kehry
                                                                                                                                                                                                                                                                                          (first entry)
                                                     95US-0431055.
94US-0234580.
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WPI; 1998-556393/47
P-PSDB; AAW71751.
cells
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Increased proliferation of B cells in the presence of membrane-bound s in culture CD40 ligand . by incubating them

Example 1; 37pp; English.

method of the invention. The method has been developed for proliferating B:cells to increase their number at least 100 fold. The method comprises: (a) providing high density, membrane bound cp40 ligand; and (b) culturing one or more B cells in the presence of this ligand. The culture results in a proliferation in the number of B cells of at least 100 fold. Also described is a method as above where the B cells are induced to differentiate into antibody-producing cells in the presence of one or more cytokines. The method can be used for stimulating B-cell proliferation in vitro or in vivo, e.g. for treating conditions in which B-cell proliferation and activation is suppressed. Eight rounds of division over six days can be achieved, corresponding to a 256-fold increase in cell numbers, which is a vast increase compared to previous

Sequence 840 BP; 263 A; 182 c; 181 G; 214 T; 0 other

Length

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                                          CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
ttcgagagaatcttactcagagctgcaaatacccacagttccgccaaaccttgcgggcaa
                                                                                      ctgaccgttaaaagacaaggactctattatatctatgcccaagtcaccttctgttccaat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                780;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98.8%;
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Pred. No. 6.4e-214;
0; Mismatches 6;
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05-DEC-1991;
23-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1525
                                                                                                                             Sequence 840
                                                                                                                                                proliferation of
                                                                                                                                                        invention relates to anti-human CD40-L monoclonal antibodies M90 secret
by hybridoma hCD40L-M90 (ATCC HB-12055) and M91 secreted by hybridoma
hCD40L-M91 (ATCC-HB-12056). M90 and M91 inhibit CD40-L binding to CD40
and the ability of trimeric CD40-L and anti-Immunoglobulin M to induce
                                                                                                                                                                                                                       Disclosure;
                                                                                                                                                                                                                                           Anti-human CD40-Ligand monoclonal antibodies
                                                                                                                                                                                                                                                               P-PSDB;
                                                                                                                                                                                                                                                                                            Spriggs.
                                                                                                                                                                                                                                                                                                                                                                           24-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                              05-OCT-1999
                                                                                                                                                                                                                                                                                                                                                                                                                US5961974-A
                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                        peripheral blood B cell; proliferation inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CD40-L;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human CD40-L
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                                    sequence encodes the human CD40 receptor ligand (CD40-L).
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atgaaaatttttatgtatttacttactgttttttcttatcacccagatgattgggtcagca 165
          ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG
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                                                                                                                                                                                                                                                                                            MK,
                                                                                       Similarity
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                                                                            Conservative
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91US-0805723.
92US-0969703.
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99.2%;
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Pred. No. 6.4e
0; Mismatches
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D40-L; anti-Immunoglobulin M;
                                                                                                                                                B cells.
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                                                                                                                                                                                                                                                                                                   RESULT
                                                                                                                          gp39; CD40 ligand; osteoblast cell death; apoptosis; be osteoporosis; osteonecrosis; inflammatory arthritis; es ovariectomy; historectomy; lipus nephritis; Takayasu's Wegener's granulomatosis; nephritis; myositis; sclerode
                                                                                        Homo
                                                                                                                                                                                         Nucleotide sequence of human gp39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CD40 ligands are used for reducing osteoblast cell death or apoptosis, and for treating or preventing bone loss in animals, preferably humans, at risk of, or undergoing, bone loss in he bone loss is associated with osteoporosis, osteoneorosis, inflammatory arthritis, post-menopausal pestrogen loss, esterogen loss due to overiore.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 879
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 113; 118pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      apoptosis
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ddiopathic autoimmune thrombocytopenia, asthma, a chronic obstructive lung disease, nephrotic/nephritic.syndrome, or cancer. They may also be used to treat or prevent bone loss in a subject undergoing, or scheduled for, an organ or bone marrow transplant.
481 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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                                         GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG
                                                                                                                                          aacaaagaggagacgaagaaagaaaacagctttgaaatgcaaaaaggtgatcagaatcct
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                          gctgaaaaaggatactacaccatgagcaacaacttggtaaccctggaaaatgggaaacag
                                                                                 caaattgcggcacatgtcataagtgaggccagcagtaaaacaacatctgtgttacagtgg
                                                                                                              CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
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                                                                                       Inducing diseases
                                                                                                                                                                                                                                                                                                                                                            cell death inhibition; stress-induced; immunosuppressive; anti-thyroid; anti-inflammatory; anti-diabetic; anti-rheumatic; anti-anaemic; opthalmological; anti-psoriatic; nephrotophic; hepatotropic; virucide;
                                                                                                                        P-PSDB; AAY96993
                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAA51745;
                                                                                                                                 WPI; 2000-452387/39
                                                                                                                                                                                                                                       06-JUL-2000
                                                                                                                                                                                                                                                                                                              Key
                                                                                                                                                                                                                                                                                                                                                                                           CD40 ligand; CD40; T cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                     31-OCT-2000
                                                                    Disclosure;
                                                                                                                                                      Newell MK,
                                                                                                                                                                         (UYVE-') UNIV VERMONT & STATE AGRIC
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                                                                                                                                                                                                                                                          WO200039283-A1.
                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                    dermatological; cytostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                Human CD40 ligand cDNA
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                                                                    Page 46;
                                                                                                                                                      Wagner D,
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                                                                                                                                                                                                                   99WO-US30930
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                                                                                       T cells with a CD40-binding agent
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CD40 engagement on T cells can be used to induce T cell receptor rearrangement and enhance T cell affinity for a particular antigen. CD engagement can be bought about by contacting CD40 with a CD40-binding agent, e.g. human CD40 ligand. The CD40-binding agents can also be used in methods for promoting T cell maturation, inhibiting T cell receptor

rearrangement,

environmental

stress-induced

cell

death

CD40

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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antigen can be inhibited, especially for a self-antigen in an autoimmune disease, which includes rheumatoid arthritis, uveitis, insulin-dependent diabetes mellitus, haemolytic anaemias, rheumatic fever, Crohn's disease Guillain-Barre syndrome, psoriasis, thyroiditis, Grave's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 altering the specificity of a T cell towards an antigen, inducing T cell reactivity towards an antigen or enhancing environmental stress-induced cell death (all claimed). T cell affinity maturation towards a specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  lupus erythematosus. Inducing environmental stress-induced T cell death is carried out in a cancerous T cell or a self-reactive T cell where the environmental stress is a chemotherapeutic agent (claimed).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   myasthenia gravis, glomerulonephritis, autoimmune hepatitis or systemic
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                             gtgactgatccaagccaagtgagccatggcactggcttcacgtcctttggcttactcaaa
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                                                         GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
                                                                                          caatccattcacttgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaat
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40

Query Match Best Local Matches

780;

Conservative

0;

Similarity

98.8%;

Score 776.4; Pred. No. 9e-2:0; Mismatches

6, 23;

Indels Length

Gaps

0

60 99

1816; 0,:

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RESULT AAS865711
ID AAS865711
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                                                                                                      cc polynucleotides are also used in diagnostics as expressed sequence tags cfor identifying expressed genes. (I) is useful in gene therapy techniques ct to restore normal activity of (II) or to treat disease states involving c(II). (II) is useful for generating antibodies against it, detecting or creat disease states involving c(II). (II) is useful for generating antibodies against it, detecting or creat cond supplement. (II) and its binding partners are useful in medical cimaging of sites expressing (II). (I) and (II) are useful for treating cC disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in cC diagnostics, forensics, gene mapping, identification of mutations cC responsible for genetic disorders or other traits to assess biodiversity cand to produce other types of data and products dependent on DNA and cc amino acid sequences. AAS64197-AAS94564 represent novel human cC diagnostic coding sequences of the invention.

CC Note: The sequence data for this patent did not appear in the printed content of the product of the complete content of the printed content of the pr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         polypeptide (II) sequences. (I) is useful as hybridisation polymerase chain reaction (PCR) primers, oligomers, and for and gene mapping, and in recombinant production of (II). The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess biodiversity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID No 22375; 103pp; English.
    Sequence 1816
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23-AUG-2000;
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                                                                                                                               secretion;
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                                                                                                                                                                                                  ctttttgctgtgtatcttcatagaaggttggacaagatagaagatgaaaggaatcttcat
                                                                                                                       gaagattttgtattcatgaaaacgatacagagatgcaacacaggagaaagatccttatcc
            NACAAAGAGGAGAAGAAAGAAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                              ttactgaactgtgaggagattaaaagccagtttgaaggctttgtgaaggatataatgtta
                                                                  TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA
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Query Match
Best Local
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07-JUN-1995;
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diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Armitage RJ, Fanslow WC, Srinivasan S;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     mutation
                                                                                                                                                                                                                                                   muteins have a higher binding affinity for human CD40 than native CD401 and can be used in binding assays.

(Note: The present sequence does not appear in the specification; it has been produced by modifying the wild-type coding sequence, which is given on pages 19-20, at codon 194).
                                                                                                                                                                                                                                                                                                                                       This sequence covers cDNA coding for human CD40L mutein polypeptides in which the wild-type Cys residue at position 194 has been replaced by a Trp, Ser, Asp or Lys residue. The muteins, designated C194W, C194S, C194D and C194W, are functional analogues of CD40L and as such are useful to induce B cell proliferation and antibody secretion. This property may be used to treat disorders associated with low levels of B cells or antibody secretion. The new CD40L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (IMMV ) IMMUNEX CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-052320/05.
P-PSDB; AAW09113, AAW09114,
                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New CD40
                                                                                                                              Local Similarity 98:
hes 777; Conservative
ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG
                                                    ATGATCGAAACATACCAAACTACTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Page -;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ligand mutein with higher CD40 affinity than native ligand in binding assays, and for therapy of disorders and involving low levels of B cells and antibody secretion
                                                                                                                                                                                                                      840 BP;
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95US-0477733
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/*tag= 1
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/note= "Encodes CD40L
position 194 if
or Lys"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31pp;
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New nucleic acid encoding human gp39 T cell antigen
ligand for the CD40 receptor, causing proliferation
differentiation of B cells and some cancer cells
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B-cell differentiation;
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Copyright (c) 1993 - 2000 Compugen Ltd.
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May 30, 2002, 05:22:46 ; Search time 8356.06 Seconds (without alignments) 2035.054 Million cell updates/sec US-08-982-272-4 786 1 ATGATCGAAACATACAACCA......TTGGCTTACTCAAACTCTGA 786 43959072 21979536 seqs, 10817449327 residues Total number of hits satisfying chosen parameters: OM nucleic - nucleic search, using sw model IDENTITY_NUC Gapop 10.0 , Gapext 1.0 Minimum DB seq length: 0 Maximum DB seq length: 200000000 Title: Perfect score: Sequence: Scoring table: Searched: Run on:

Database :

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Description	Sequence 4, Appli	3, A	e 1,	7,7	Ä	Ļ	Sequence 10, Appl	11,		7, A	٦,	Sequence 1, Appli	'n	Sequence 1, Appli	11,	Sequence 11, Appl	Sequence 7, Appli	Sequence 1, Appli	11,	Sequence 11, Appl	1, A	1,	٦,	11,	11,	1, 2	Sequence 7, Appli				
ID	US-08-982-272-4	US-08-107-353-3	US-08-982-272-1	PCT-US94-00786-7	PCT-US94-14767-1	PCT-US97-11956-1	US-07-805-723-10	US-07-969-703A-11	US-07-969-703B-11	US-08-009-258-7	\sim	US-08-215-862-1	US-08-234-580-3	US-08-396-230-1	US-08-477-733A-11	US-08-484-624-11	US-08-589-771-7	US-08-677-762-1	US-08-770-981-11	-60-	US-09-322-021-11	US-09-322-021A-11	-365-940-	US-09-365-940A-11	US-09-387-935-1	US-09-388-079-1	-09-388-079	US-09-392-618-11	US-09-399-106-11	-08-430-4	US-09-545-993-7
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Mismatches

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			, 8 8 8 8 9 8 8 8 8	9.66		NESCULA NS-08-982-272-4 Sequence 4, Application 1 GENERAL INFORMATION: APPLICANT: Kipps, Th	CANT: Sharma, S. CANT: Cantwell, OF INVENTION: OF INVENTION:	TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION:	: H & '' ?	KEET: Suite 4700 Y: Los Angeles ATE: California JNTRY: U.S.A.	ZIP: 900/1-2000 COMPUTER READABLE FORM MEDIUM TYPE: 3.5" D	COMPUTER: 15M COMPACT. OPERATING SYSTEM: 15M SOFTWARE: FASTSEQ VER CURRENT APPLICATION DATA	APPLICATION NUMBER: CO. ELING DATE: CLASSIFICATION: PRIOR APPLICATION DATA:	APPLICATION NUMBER: 00/1 FILING DATE: 12/9/96 ATTORNEY/AGENT INFORMATION: NAME: Guise, Jeffrey W.			S. K.	TOPOLOGY: linear 982-272-4	Query Match Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 786;
                                                  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/107,353
FILING DATE: 19930813
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 98.8%; Score 776.4; DB 5; Best Local Similarity 99.2%; Pred. No. 7.3e-197; Matches 780; Conservative 0; Mismatches 6;
                                              PC-DOS/MS-DOS
                                                                                                                                            NAME: PETKINS, PATTICIA A
REGISTRATION NUMBER: 34,693
REFRENCE/COCKET NUMBER: 100.
TELECOMMUNICATION INFORMATION:
TELEPAN: (206)233-0644
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                             34,693
                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: CDNA to mRNA
                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                          1: 786 base pairs
nucleic acid
                                                                                                                                                                                                                                                                                                        linear
                                                                                                                                                                                                                                                                                                                                                                                                                                     1..783
                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Human
STRAIN: CD40-L
                                                                                                   FILING DATE: 19 CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                  ANTI-SENSE: NO
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY:
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US-08-107-353-3
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                                                                                                                                                                                                                                                             LENGTH:
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                                                                                                                                                        TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NOVEL EXPRESSION VECTORS
CONTAINING ACCESSORY
MOLECULE LIGAND GENES AND
THEIR USE FOR IMMUNO-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: 1.44 Mb storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSeq Version 2.0
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/08/982,272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/08982272 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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ATTORNEY/AGENT INFORMATION:
NAME: Glise, Jeffrey W.
REGISTRATION NUMBER: 34,613
REPRENCE/POCKET NUMBER: 231/1
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Kipps, Thomas J. APPLICANT: Sharma, Sanjai APPLICANT: Cantwell, Mark
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TELEX: 67-3510
INFORMATION FOR SEQ. ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 786 base pairs
TYPE: nucleic acid
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MODIFIED OF INVENTION:
MODIFIED OF INVENTION:
MODIFIED OF INVENTION:
MODIFIED OF SEQUENCES:
CORRESPONDENCE ADDRESS:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 6
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STATE: California
COUNTRY: U.S.A.
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781 CTCTGA 786
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US-08-982-272-1
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DETECTION AND TREATMENT OF MUTATIONS

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TITLE OF INVENTION:
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                                                                                                                                                                                                                        61 AIGAAAAITITTATGIAITTACITACTGITITCCTTAICACCCAAAIGAITGGAICTGIG 120
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                                          Gaps
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                                                            1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
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                        DB 13; Length 786;
                                  7.3e-197
                                            0; Mismatches
                           Score 776.4;
                                    Pred. No.
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GENERAL INFORMATION:
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WILLIAM
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MELANIE
                           98.8%;
99.2%;
                          Query Match
Best Local Similarity 99.29
Matches 780; Conservative
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RENSHAW,
SPRIGGS,
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781 CTCTGA 786
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PCT-US94-00786-7
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APPLICANT:
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 US-08-982-272-1
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181 GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC 240
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99.2%; Pred., No. 7.5e-197;
Live 0; Mismatches 6;
                                                                                                                                                              SOFTWARE: MS MORE for Apple 5.1, Version a CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/00786
FILING DATE:
                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/009,258
APPLING DATE: 01/22/93
ATTORNEY/AGENT INFORMATION:
NAME: PERKINS, PATRICIA ANNE
REGISTRATION NUMBER: 34,693
                                          SI IMMUNEX CORPORATION STREET
                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
IN A CD40
                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple OF
                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 99.2
Matches 780; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                        2065870606
                               CORRESPONDENCE ADDRESS
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                   NUMBER OF SEQUENCES:
                                                                                          WASHINGTON
                                                                                                                                                                                                                                                           CLASSIFICATION:
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CLONE: CD40-L
                                                          STREET: 51 UNACTIVE SEATTLE
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ORIGINAL SOURCE:
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                                                                                                             USA
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, LOCATION:
PCT-US94-00786-7
                                                                                                                           ZIP: 98101
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                                                 ADDRESSEE:
                                                                                                          COUNTRY:
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                                                                                             STATE:
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                                                                                                                                                          466 GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG 525
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AACAAAGAGGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 360
                                                                                                                                                                                                                        526 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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                CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAAACAACATCTGTGTTACAGTGG
                                                                                                                                         APPLICANT: LONGO, DAN L.
APPLICANT: MURPHY, WILLIAM
TITLE OF INVENTION: METHOD OF PREVENTING OR TREATING
TITLE OF INVENTION: DISEASE CHARACTERIZED BY NEOPLASTIC CELLS
TITLE OF INVENTION: EXPRESSING CD40
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: Apple Macintosh System 7.1
SOFTWARE: Microsoft Word for Macintosh, Version #5.1a
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USSN 08/172,664
er 23, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: PCT/US94/14767
FILING DATE: December 21, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Application PC/TUS9414767 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              3: IMMUNEX CORPORATION 51 UNIVERSITY STREET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: ARMITAGE, RICHARD APPLICANT: FANSLOW, WILLIAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN
FILING DATE: December 23,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: WASHINGTON
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| 826 CTCTGA 831
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CITY: SE
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661 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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Pred. No. 7.5e-197;
); Mismatches 6;
FELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                           98.8%;
            TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                 IMMEDIATE SOURCE:
CLONE: CDAA-*
                                                              SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                        46..831
                                                                                                                                        linear
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                                                                                                                                     TOPOLOGY: line
MOLECULE TYPE: HYPOTHETICAL: N
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ORGANISM: HOM
                                                                                                                                                                                                                                                                                                      NAME/KEY:
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Matches 780;
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360 405 420 465 480 525 645

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721 GIGACIGAICCAAGCCAAGIGAGCCAIGGCACIGGCIICACGICCIIIGGCIIACICAAA 780,
                Length 840;
                                                                                                                                                                                   Sequence 1, Application PC/TUS9711956
GENERAL INFORMATION:
APPLICANT: Immunex Corporation
TITLE OF INVENTION: METHOD OF ACTIVATING DENDRITIC CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                   SYSTEM: Apple Operating System 7.6 Microsoft Word for Macintosh, Version #6.0.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATE: USBN 08/763,995 APPLICATION NUMBER: 12-DEC-1996 ATTORNEY/ACBNT INFORMATION: NAME: PERKINS, PARTICIA A REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USSN 08/677,762
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCT/US97/11956
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MEDIUM TYPE: Floppy disk
COMPUTER: Power Macintosh 7200/90
                                                                                                                                                                                                                                                                                           E: IMMUNEX CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REGISTRATION NUMBER: 34,69:
REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 10-JUL-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                                                                WASHINGTON
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                                                                           CTCTGA 786
                                                                                             TELEPHONE:
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LOCATION:
PCT-US97-11956-1
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                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                          PCT-US97-11956-1
                                                                                                                                                                                                                                                                                                                                   CITY:
STATE:
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GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG 480 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT 540 526 CIGACCGITAAAAGACAAGAACICIATIATATATGIGAGCCAAGICACCITCIGIIIIIIIII

466 481

421

361 CAAATTGCGGCCACATGTCATAAGTGAGGCCAGCAGTAAAAACAACATCTGTGTTACAGTGG

301 AACAAAGAGGAGAAGAAAAAAAAAAAGTTTGAAATGCAAAAAGGTGATCAGAATCCT

CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT

106 121

ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120 241 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300

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721 GIGACIGAICCAAGCCAAGIGAGCCAIGGCACIGGCIICACGICCIIIGGCIIACICAAA 780
                                                                                                                                                                                                601 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGGAA 660
                                                                                                                   661 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                                                                      NOVEL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPPRATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IMMUNEX CORPORATION
                                                                                                                                                                                                                                                                                                                                                          Sequence 10, Application US/07805723 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                           FANSLOW, WILLIAM
SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WASHINGTON
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1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60

Score 776.4; DB 1; Pred. No. 7.5e-197; 0; Mismatches 6;

Conservative

Query Match Best Local Similarity Matches 780; Conserv

Gaps

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CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
                                                                                                                                           Gaps
                                                                        1 ATGATCGAAACATACAACCATCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
                                                                                                                            GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFWMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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Pred. No. 7.5e-197;
); Mismatches 6;
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                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: NOVEL CYTOKINE NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                      Sequence 11, Application US/07969703A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      E: IMMUNEX CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 286
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                      APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
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IMMEDIATE SOURCE:
CLONE: CD40-L
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Best Local Similarity 99.2°
Matches 780; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US
FILING DATE: 19921023
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                                                                                                                                                                                               781 CTCTGA 786
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; LOCATION:
US-07-969-703A-11
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US-07-969-703A-11
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PatentIn Release #1.0, Version #1.25
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Pred. No. 7.5e-197;
0; Mismatches 6;
                                   APPLICATION NUMBER: US/07/805,723
                                                                                                                                         REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: OSTER, JEFFREY B.
REGISTRATION NUMBER: 32585
                                                                                                                                                                                                         INFORMATION FOR SECTION 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 98.8%;
Best Local Similarity 99.2%;
Matches 780; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                                                                                                                                                              840 base pairs
                  CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: CDNA HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                      linear
                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                           ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                 STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                          ANTI-SENSE: NO
                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                      TOPOLOGY:
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US-07-805-723-10
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õ a ò a à IBM PC compatible SYSTEM: PC-DOS/MS-DOS Patentin Release #1.0, Version #1.25

OPERATING SYSTEM:

SOFTWARE: Patentin Release #1.0, Ve CURRENT APPLICATION DATA: APPLICATION NUMBER: US/07/969,703B

19921023

34,693

ATTORNEY/AGENT INFORMATION: NAME: Perkins, Patricia A. REFERENCE/DOCKET NUMBER:

CLASSIFICATION:

FILING DATE:

NAME: Perkins, Patric REGISTRATION NUMBER:

FELECOMMUNICATION INFORMATION

2065870430

TELEPHONE:

TELEFAX:

2065870606

LENGTH: 840 base pairs TYPE: NUCLEIC ACID

linear

STRANDEDNESS:

<u>Q</u>

MOLECULE TYPE: HYPOTHETICAL:

TOPOLOGY:

INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:

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US-07-969-703B-11
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                                                                             tactgaactgtgaggagattaaaagccagttgaaggctttgtgaaggatataatgtta 300.
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                                  46 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 105
                                                               121 CTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
                                                                                                              181 GAAGATTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC 240
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APPLICANT: ARMITAGE, RICHARD APPLICANT: FANSLOW, WILLIAM APPLICANT: SPRIGGS, MELANIE
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ORGANISM: Homo sapiens IMMEDIATE SOURCE:

ORIGINAL SOURCE:

ANTI-SENSE: NO

46..831

NAME/KEY:

CLONE:

Length 840;

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CIGACCGITAAAAGACAAGGACTCTATTATATTGTGCCCAAGTCACCTTCTGTTCCAAT 540
                                                                                                      106 AIGAAAATTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 165
                                                                                        ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
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                                            1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
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                        Indels
Score 776.4; DB 3;
Pred. No. 7.5e-197;
0; Mismatches 6;
   98.88;
   Query Match
Best Local Similarity 99.2
Matches 780; Conservative
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NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:

INDIVERSITY STREET

ADDRESSEE:

WASHINGTON SEATTLE USA COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk

98101

COUNTRY:

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Query Match
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                                                                                                                        CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
                                                                                                                                        GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780
               CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 600
                                                            TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 660
                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: RANITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: WIDHARI
APPLICANT: WIDHARI
TITLE OF INVENTION: IN A CD40 LIGAND GENE
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/009,258 FILING DATE: 19930122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS.
                                                                                                                                                                                                                                                                                                                                                     Sequence 7, Application US/08009258 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: PERKINS, PATRICIA A.
REGISTRATION NUMBER: 34,693
REFRENCE/DOCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 840 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear MOLECULE TYPE: CDNA
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ORGANISM: HOM
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US-08-009-258-7
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                                                                                                       61 ATGAAAATTTTTATGTATTTACTTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
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                                                                                  1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
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DISEASE CHARACTERIZED BY NEOPLASTIC CELLS
EXPRESSING CD40
                                             ö
    Length 840;
                                             Indels
  Score 776.4; DB 4;
Pred. No. 7.5e-197;
); Mismatches 6;
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Sequence 1, Application US/08172664
GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
98.8%;
99.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: LONGO, DAN L.
APPLICANT: MURPHY, WILLIAM
TITLE OF INVENTION: METHOD
TITLE OF INVENTION: EXPRESS
TITLE OF INVENTION: EXPRESS
                                           Conservative
                        Similarity
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                        Best Local Sin
Matches 780;
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us-08-982-272-4.rnpm

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406 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 465
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: SEATTLE STATE: WASHINGTON
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                                                                                                                                                                                                                                                                                                                                                                                                781 CTCTGA 786
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|826 CTCTGA 831
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COPOLOGY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-215-862-1
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                                                                                                                                                                                                Version #5.1a
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Pred. No. 7.5e-197;
0; Mismatches 6;
                                                                                                                                                                         OPERATING SYSTEM: Apple Macintosh System 7.1 SOFTWARE: Microsoft Word for Macintosh, Versicorremy Application Data: APPLICATION NUMBER: US/08/172,664 FILING DATE: December 23, 1993
                    NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS: 1 IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 34,693
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2818
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Match 98.8%;
Local Similarity 99.2%;
les 780; Conservative
                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: Apple
                                                                                                                                                                                                                                                             CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CDS
46..831
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                                                                                              WASHINGTON
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CLONE: CD40-L
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                                                                                SEATTLE
                                                                                                             USA
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, LOCATION:
US-08-172-664-1
                                                                                                                           98101
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                                                                                                              COUNTRY:
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                                                                                                STATE:
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661 CAATCCATTCACTTGGGAGGAGTATTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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                                                                                                                                                                                                                                   601 TICGAGAGAATCITACICAGAGCIGCAAAIACCCACACATICCGCCAAACCIIGCGGGCAA 660
                                                                                                                                                                                                                                                                                                                                                                                                          421 GCTGAAAAAGGATACTACACCATGAGCAACATGGTAACCTGGAAAATGGGAAAAAG 480
                                                                                                                                                                                                                                                       481 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                  541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                                                                                                                                                                  APPLICANT: Gruss, Hans-J_rgen
TITLE OF INVENTION: Method of Diagnosing or Treating Hodgkin's
TITLE OF INVENTION: Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
OPERATING SYSTEM: Apple Macintosh System 7.1
OPERATING SYSTEM: Apple Macintosh, Version #5.1a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/215,862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/08215862
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              relecommunication information Telephone: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Particia A.
RECISTRATION NUMBER: 34,65
REFERENCE/DOCKET NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (206)233-0644
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COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY
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ATTORNEY AGENT INFORMATION:
NAME: MILLMAN, ROBERT A
REGISTRATION NUMBER: 36,217
                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                    Query Match 98.8%;
Best Local Similarity 99.2%;
Matches 780; Conservative
                                                                                                                                                                                                                                       TELEPHONE: (202)371-2600
TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                             LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                  22..807
                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                      ; NAME/KEY: CDS
; LOCATION: 22...
US-08-234-580-3
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                                                                         840;
                                                                         Length
                                                                         776.4; DB 6;
No. 7.5e-197;
                                                                         Score 776.4;
Pred. No. 7.5e
0; Mismatches
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                                                                        Query Match 98.8%;
Best Local Similarity 99.2%;
Matches 780; Conservative
                                    46..831
IMMEDIATE SOURCE:
        CLONE: CD40-L
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTCTGA 831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       781 CTCTGA 786
                         ; NAME/KEY:
; LOCATION:
US-08-215-862-1
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APPLICANT: CASTLE, BRIAN E
APPLICANT: CASTLE, BRIAN E
TITLE OF INVENTION: METHODS FOR PROLIFERATING AND
TITLE OF INVENTION: DIFFERENTIATING B CELLS, AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             840;
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                                                                                         CORRESPONDED

ADDRESSEE: STERN, ALLOL

STREET: 100 NEW YORK AVE. N.W. SULL.

CITY: WASHINGTON

STATE: D.C.

ZIP: 2006.

ZIP: 2006.

COMPUTER READABLE FORM:

MEDIUM TYPE: Flopy disk

COMPUTER: IBP CC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/234,580

TTING DATE: 28+APR-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 776.4; DB 6;
Pred. No. 7.5e-197;
0; Mismatches 6;
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IMMEDIATE SOURCE:
CLONE: CD40L
                               ; NAME/KEY:
; LOCATION:
US-08-396-230-1
                        FEATURE:
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  721 GIGACTGAICCAAGCCAAGIGAGCCAIGGCACIGGCTICACGICCITIGGCTIACTCAAA 780
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                                                 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: MALISZEWSKI, CHARLES R. TITLE OF INVERTION: METHOD POR STIMULATING A CELL-MEDIATED TITLE OF INVERTION: IMMUNE RESPONSE WITH A CD40 BINDING TITLE OF INVENTION: PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: Apple Power Macintosh System 7.5.5 SOFTWARE: Microsoft Word for Macintosh, Version 6.0.1 CURRENT APPLICATION DATA:
FILLING DATE: March 1, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 34,693
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2830
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)283-0644
                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application US/08396230 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: CAMPBELL, KIM A. APPLICANT: KENNEDY, MARY K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk COMPUTER: Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
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CITY: SEATTLE
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                                                                                               CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
                                                                                                                                                                226 GAAGATTTGTATTCATGAAACGATACAGAGATGCAACAGGAGAAAGATCCTTATCC 285
                                                           46 ATGATCGAAACATACAACATCTCCCCGATCTGCGGCCACTGGACTGCCATCAGC 105
                        Gaps
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                                               ATGATCGAAACATACAAACCTTCTCCCCGATCTGCGGCCACTGGACTGCCATCAGC
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Length. 840;
                         Indels
Score 776.4; DB 7;
Pred. No. 7.5e-197;
                           9
                         0; Mismatches
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US-08-477-733A-11
Sequence 11, Application US/08477733A
GENERAL INFORMATION:
     98.8%;
99.2%;
                             Conservative
    Query Match
Best Local Similarity
Matches 780; Conserv
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OPERATION APPLE MACHICOM
OPERATION SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, version 5.1a
CURRENT APPLICATION DATA:
                                                                                                 NOVEL CYTOKINE THAT BINDS CD40
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/477,733A
                   APPLICANT: FANSLOW, WILLIAM
APPLICANT: SFRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2802-D
                                                                                                                                                            E: IMMUNEX CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
PHOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08/249,189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRICE APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEPAX: 2065870606
ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/24
FILING DATE: May 24, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
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TYPE: nucleic acid
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IMMEDIATE SOURCE:
CLONE: CD40-L
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
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MOLECULE TYPE: CDNA
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                                                                                                                                                                                                    SEATTLE
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US-08-477-733A-11
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481 CTGACCGTTAAAAGACAAAGGACTCTATTATATCTATGCCCCAAGTCACCTTCTGTTCCAAT 540
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                                                                                                                 GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC
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Job time: 18652 sec
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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May 30, 2002, 05:38:54 ; Search time 627.19 Seconds (without alignments) 1315.293 Million cell updates/sec US-08-982-272-4 786 1 ATGATCGAAACATACAACCA......TTGGCTTACTCAAACTCTGA 786 Scoring table: Perfect score: Sequence: е :: Run

IDENTITY_NUC Gapop 10.0 , Gapext 1.0

1701006 Total number of hits satisfying chosen parameters: 850503 segs, 524770293 residues Searched:

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries Minimum DB seq length: 0 Maximum DB seq length: 200000000

/cgn2_6/ptodata/1/pna/US07_NEW_COMB.seq:* /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq:* /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq:* /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq:* /cgn2_6/ptodata/1/pna/US60_NEW_COMB.seq:* Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Gaps

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9:

0; Mismatches

Matches 780; Conservative Query Match Best Local Similarity

Length 1816; Indels

Score 776.4; DB 5; Pred. No. 2.1e-193;

98.8%; 99.2%;

ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120 CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180 GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC 240 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300

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CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAAACAACATCTGTGTTACAGTGG 420

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Description				131,	Sequence 9, Appli	203		664,	7000,			20614			1996	239,	1323		Sequence 83795		824;	27253	Sequence 514, App	11996	Sequence 241769,	
ID	US-09-053-375B-226	US-09-442-384B-457	US-09-053-375B-821	US-09-911-904-131	US-09-875-453A-9	PCT-US02-10421-202	US-10-112-699-202	US-09-919-002-664	US-09-540-210B-7000	US-09-540-210B-14649	US-10-123-155-120	US-10-027-632-206146	US-10-027-632-206147	US-09-539-331D-26090	US-10-105-299-1996	US-10-106-698-239	US-09-789-189-1323	US-10-027-632-83794	US-10-027-632-83795	US-09-442-384B-628	US-10-105-299-824	US-10-027-632-27253	US-09-053-375B-514	US-09-919-002-11996	US-10-027-632-241769	050130 000 500 01 011
å Query Match Length DB	1816 5	1816 5	1250 5	508 5	2395 5	569 1	9 699	2127 5	234 5	230 5	775 6	9 699	9 699	438 5	1751 6	1751 6	418 5	621 6	621 6	10351 5	803 6	766 6	2259 5	646 5	695 6	2 3031
& Query Match	98.8	98.8	72.7	45.9	18.8	5.8	5.8	5.6	5.0	4.9	4.9	4.8	4.8	4.7	4.6	4.6	4.6	4.6	4.6	4.6	4.5	4.5	4.5	4.5	4.5	7
Score	776.4	776.4	571.6	360.6	147.8	45.6	45.6	44	39.4	38.2	38.2	38	38	36.8	36.4	36.4	35.8	35.8	Ŋ.	35.8	35.6	35.4	35.4	35.2	35.2	30
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S	FILE REFERENCE: CLON-006 CURRENT APPLICATION NUMB. CURRENT FILING DATE: 19 NUMBER OF SEQ ID NOS: 15 SOFTWARE: FastSEQ for W1 SEQ ID NO 226 LENGTH: 1816 LENGTH: 1816 TYPE: DNA ORGANISM: HOMO Sapiens US-09-053-375B-226	ILE REFERENCE: CLON URRENT APPLICATION URRENT FILING DATE: UMBER OF SEQ ID NOS OF TOWARE: FASTSEQ fO OF TOWARE: 1816 LENGTH: 1816 LENGTH: 1816 LENGTH: 1816 OF TOWARE: PASTSED OF	XENCE: CL PLICATIO LING DAT SEQ ID N FastSEQ 226 816 A HOMO SA	CLON-006 ION NUMBER: US/0 ATE: 1998 08-31 NOS: 1543 Q for Windows Ve		FILE REFERENCE: CLON-006 CUGRENT APPLICATION NUMBER: US/09/053,375B CURRENT FILING DATE: 1998-08-31 NUMBER OF SEQ ID NOS: 1543-08-31 NUMBER OF SEQ ID NOS: 1543-08-31 SOFTWARE: FastSEQ for Windows Version 4.0 EQ ID NO 226 LENGTH: 1816 TYPE: DNA ORGANISM: Homo sapiens 09-053-375B-226		

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TYPE: DNA
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Pred. No. 2.1e-193;
0; Mismatches 6;
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TITLE OF INVENTION: Hematology/Immunology Array
FILE REFERENCE: CLON-006GIP15
FULE REFERENCE: CLON-006GIP15
CURRENT FILING DATE: 1999-11-17
PRIOR APPLICATION NUMBER: 09/053,375
PRIOR APPLICATION NUMBER: 09/053,375
PRIOR FILING DATE: 1998-03-31
NUMBER OF SEQ ID NOS: 830
SCFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 457
LENGTH: 1816
                                                                                                                                                                                                                                                                                                                 Sequence 457, Application US/09442384B
GENERAL INFORMATION:
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Best Local Similarity 99.2%;
Matches 780; Conservative
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US-09-442-384B-457
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AACAAAGAGGAGACGAAGAAAGAAAAGAAACAGCTTTGAAAATGCAAAAAGGTGATCAGAATCCT
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Pred. No. 6.6e-140;
0; Mismatches 124;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Bibilashvilli, Robert
TITLE OF INVENTION: Nucleic Acid Arrays
FILE REFERENCE: CLON-006
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GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
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83.8%;
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Tam, Albert W.
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Velligan, Mark D.
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Thomas, Rita L.
Kongpachith, Ana
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                                      gaagattttgtattcataaaaagactaaagagatgcaacaaaggagagaggatctttatcc
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.7e-85;
es 84;
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Pred. No. 7.7e-
0; Mismatches
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APPLICANT: Pickett, Gavin G.
APPLICANT: Neft, Robin Eileen
APPLICANT: Dunn, II, Robert Thomas
TITLE OF INVENTION: CANINE TOXICITY GENES
FILE REFERENCE: 400742000200
CURRENT APPLICATION NUMBER: US/09/911,904
PRIOR APPLICATION NUMBER: US 60/220,057
PRIOR FILING DATE: 2000-07-21
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SOFWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 131
LENCTH: 508
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83.0%;
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US-09-911-904-131
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Best Local Similarity 83.0
Matches 424; Conservative
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CAGCTTTGAAATGCAAAAAGGTGATCAGAATCCTCAAATTGCGGCACATGTCATAAGTGA
                                                                                                                    GGCCAGCAGTAAAACAACATCTGTGTTACAGTGGGCTGAAAAAGGATACTACACCATGAG
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TITLE OF INVENTION: PROMOTERS FOR REGULATED GENE EXPRESSION
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Pred. No. 5.4e
0; Mismatches
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CURRENT APPLICATION NUMBER: US/09/875,453A
CURRENT FILING DATE: 2002-05-06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR FILING DATE: 2000-06-06
NUMBER OF SEQ ID NOS: 246
SOFTWARE: FastSEQ for Windows Version 4.0
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88.0%;
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Best Local Similarity 88.0
Matches 161; Conservative
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ID NOS: 2959

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SEQ ID NO
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF CANCER
FILE REFERENCE: 210121.565
CURRENT FILLING NUMBER """
                                                                                                                                                                                                                                                                                                                                                    TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIAGNOSIS OF CANCER
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0; Mismatches 119;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 45.6;
                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION NUMBER: PCT/USO2/10421
CURRENT FILING DATE: 2002-03-28
NUMBER OF SEQ ID NOS: 2959
                                                                                                                                                                                                                                                         Application PC/TUS0210421
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                                                                                                                                                                                                                                                                                                                              Bangur, Chaitanya S.
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Matches 117; Conservative
                                                                                                                                                                                                                                                                                     APPLICANT: Corixa Corporation
                                                                                                                                                                                                                                                                                                                                           Gaiger, Alexander
                                                                                                                                                                                                                                                                                                                                                                                 FILE REFERENCE: 210121.565PC
                                                                                                                                                                                                                                                                                                   Wang, Tongtong-Wang, Siqing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
CORGANISM: Homo sapiens
PCT-US02-10421-202
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Pred. No. 0.072;
0; Mismatches 120;
                                                                                                                                                 0; Mismatches 119;
                                                                                                   Score 45.6; DB
Pred. No. 0.019;
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GENERAL INFORMATION:
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                                                                                                                                                        Conservative
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Best Local Similarity
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Best Local Similarity
Matches 117; Conserv
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TYPE: DNA
CRGANISM: HOMO
US-10-112-699-202
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LENGTH: 2127
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Query Match 5.0%; Score 39.4; DB 5; Length 2: Best Local Similarity 48.8%; Pred. No. 0.61; Matches 103; Conservative 0; Mismatches 108; Indels
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GENERAL INPORMATION:
APPLICANT: Sellhamer, Jeffrey J.
APPLICANT: Delegeane, Angelo M. (APPLICANT: Stuart, Susan G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-540-210B-14649
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ANT: Naughton, Rebecca E.
OF INVENTION: POLYNUCLEOTIDES OF URINARY TRACT TISSUE
REFERENCE: PD-1037 CIP
                                                                     RRENT APPLICATION NUMBER: US/09/540,210E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           June 25, 1996
NUMBER: 08/903,802
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TON NUMBER: 60/025,204
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383 GTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGGGCTGAAAAAGGATACTACACCA 442

443 TGAGCAACACTTGGTAACCCTGGAAAATGG 473

263 AAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTAAACAAAGAGGAGACGAAGAAAG 322

Length 234;

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; LOCATION: 24, 113
; OTHER INFORMATION: a, t, c, g, or other
US-09-540-210B-7000
                                                                                                                                                                                                                                       LING DATE: May 29, 1997
PLICATION NUMBER: 09/107,592
                                                                                                                                                 PLICATION NUMBER: 09/050,817
                                                                                                                                                                                                                 :LING DATE: May 8, 1998
PLICATION NUMBER: 60/048,431
                                      ILING DATE: February 13, 1997
                                                                                                                                                               LING DATE: March 30,
                                                                                                                                                                                                                                                                 June 30,
                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo sapiens
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SOFTWARE: PERL PR
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LENGTH: 234
PRIOR
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APPLICANT: Naughton, Rebecca E.
FITLE OF INVENTION: POLYNUCLEOTIDES OF URINARY TRACT TISSUE
FILE REFERENCE: PD-1037 CIP
                                                             CURRENT APPLICATION NUMBER: US/09/540,210B
CURRENT FILING DATE: 2002-04-03
PRIOR APPLICATION NUMBER: 08/972,899
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                                                                                                                                    FILING DATE: November 18, 1997
APPLICATION NUMBER: 08/395,244
FILING DATE: February 27, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         60/023,308
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NG DATE: April 10, 1996
ICATION NUMBER: 08/755,
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INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 230;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No: hu00678829
US-09-540-210B-14649
                                  APPLICATION NUMBER: 09/050,817
FILING DATE: March 30, 1998
APPLICATION NUMBER: 60/043,792
FILING DATE: April 11, 1997
APPLICATION NUMBER: 09/074,999
                                                                                                                                                                                                                                        DATE: July 1, 1997
ATION NUMBER: 09/094,079
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APPLICATION NUMBER: 60/040,199
                                                                                                                                                  60/048,431
                                                                                                                                                                     G DATE: MAY 29, 1997
CATION NUMBER: 09/107,592
                                                                                                                                                                                                                       CATION NUMBER: 60/052,751
                    (March 14, 1997
                                                                                                                                                                                                  June 30, 1998
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                                                                                                                                                                                                                                                                              PRIOR FILING DATE: June 9, 1
PRIOR APPLICATION NUMBER: 60
PRIOR FILING DATE: June 13,
NUMBER OF SEQ ID NOS: 35654
SOFTWARE: PERL PROGRAM
SEQ ID NO 14649
LENGTH: 230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 50.8
Matches 91; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Watanabe, Colin
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                                                                                                                                   LING DATE: May 8,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           umas, Danie
                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens
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Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ERLGPRPMLEVIEDCGGWDLGGAEERPGVAARWDLNRLLYKAQGVYSAAALFSLTVSLDD 253
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TCAGAATCCTCAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGT 410
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  111 TGGATCTGTGCTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAG 170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 206146, Application US/10027632
GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
                                                                                                                                                                                                                                                                                                                                                                             Length 775,
                                                                                                                                                                                                                                                                                                                                                                                                                                      45; Conservative 192; Mismatches 404; Indels
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                                                                                                                 See Palm or File Wrapper
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9
  ENCODING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                Score 38.2;
                                                     CURRENT APPLICATION NUMBER: US/10/123,155 CURRENT FILING DATE: 2002-04-15
                                                                                                                                                                                                                                                                                                                                                                                                              Pred. No. 1
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                                                                                                                                                                                                                                                                                                                                                                                                              . 08;
                       FILE REFERENCE: P3330R1C30
                                                                                                        Prior Application removed NUMBER OF SEQ ID NOS: 550 SEQ ID NO 120
                                                                                                                                                                                                                             ; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-123-155-120
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Best Local Similarity
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TITLE OF INVENTION:
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                                                                                                                                                                                                  LENGTH: 775
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176 TTCATGAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGAGAAAGATCCT 235
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293 gagatatcagggagctgaaaaatacaatgaataatgtgaaaatgcaatagagag 346
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9
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PRIOR PELICATION NUMBER: US 60/198,676
PRIOR PELICATION NUMBER: US 60/198,676
PRIOR PELICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-02-24
PRIOR PELICATION NUMBER: US 60/165,218
PRIOR PELICATION NUMBER: US 60/167,363
PRIOR PELICATION NUMBER: US 60/167,363
PRIOR PELICATION NUMBER: US 60/156,358
PRIOR PELICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR PILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-24
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
                                                                                                                                                                                                                                                                                                                          PRIOR FILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
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SOFTWARE: FastSEQ for Windows Version 4.0
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Pred. No.
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Best Local Similarity
Matches 89; Conserv
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US-10-027-632-206146
                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 206146
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Length 663

DB 6;

Score 38;

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Search completed: May 30, 2002, 05:39:00
Job time: 16714 sec
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                                                                                                                                                                                                                                                               236 TATCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAA 295
                                                                                                                                                                                                                                                                                                                                  233 aacagtaagtgaccagaatgagaaatttaacagagattgaaataatttttaagaatcaaa 292.
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                                                                                                    176 ITCATGAAGATITIGIATICATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCT 235
                                                                                                                                                                               cagtatgcagagggaagatgagaggggctnccccangaaagaagactctggtctnca 290
                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OF INVENTION: POLYNUCLEOTIDES OF CARDIOVASCULAR SYSTEM TISSUE REFERENCE: PD-1022 CIP
                                                                                                                                                                                                                                                                                                                                                                                                                          296 TGTTAAACAAAGAGGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTG 349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     293 gagatatcagggagctgaaaaatacaatgaataatgtgaaaatgcaatagagag 346
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                                     Indels.
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Pred. No. 1.9;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No: hu00341082
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LOCATION: 262, 268, 288, 330, 338, 353, 371

COCATION: 26, 268, 269, 330, 358, 353, 371

SCOPPER INFORMATION: a, t, c, g, or other

COCATION: a, t, c, g, or other

SCOPPER STATE ST
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TITLE OF INVENTION: Human Secreted Proteins
FILE REFERENCE: PS950
CURRENT APPLICATION NUMBER: US/10/105,299
CURRENT FILING DATE: 2002-03-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LICATION NUMBER: US/09/539,331D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 26090, Application US/09539331D GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1996, Application US/10105299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Seilhamer, Jeffrey J.
Delegeane, Angelo M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2000-03-30
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illarity 49.7%;
Conservative
Best Local Similarity 51.1%;
Matches 89; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prior Application removed NUMBER OF SEQ ID NOS: 409
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
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US-09-539-331D-26090
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Matches 89,
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1174 TAACTGAAGTGTGAAATAACAAAAGAATGCATATAAGTCCGTTATTAAAAATCCCTGTTA 1115
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                                                                                                                                                                                                                                                                                                                        Length 1751;
                                                                                                                                                                                                                                                                                                                                                                      66; Indels
NUMBER OF SEQ ID NOS: 15197
Prior Application removed - See File Wrapper or Palm
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                  9
                                                                                                                                                                                                                                                                                                                                  DB
                                                                                                                                                                                                                                                                                                                             Score 36.4; DE Pred. No. 6.6; 0; Mismatches
                                                                                                                                                                                              ; NAME/KEY: misc_feature
; LOCATION: (1741) .(1742)
; OTHER INFORMATION: n equals a,t,g, or
US-10-105-299-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1054 GAATTGGTGTAAATGTTTGA 1033
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            361 CAAATTGCGGCACATGTCATAA 382
                                                                                                                                                                                                                                                                                                                               Query Match 4.6%;
Best Local Similarity 53.5%;
Matches 76; Conservative
                                                                                                                                                       ORGANISM: Homo sapiens
                                                                                SEQ ID NO 1996
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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OM nucleic - nucleic search, using sw model

May 30, 2002, 02:51:25 ; Search time 139.75 Seconds (without alignments) 1381.524 Million cell updates/sec Run on:

US-08-982-272-4

786 1 ATGATCGAAACATACAACCA.....TTGGCTTACTCAAACTCTGA 786 Title: Perfect score: Sequence:

Scoring table:

IDENTITY_NUC Gapox 10.0

Total number of hits satisfying chosen parameters:

383533 seqs, 122816752 residues

Searched:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*/ /cgn2_6/ptodata/2/ina/backfiles1.seq:* /cgn2_6/ptodata/2/ina/5B_COMB.seq:*/cgn2_6/ptodata/2/ina/6A_COMB.seq:*/cgn2_6/ptodata/2/ina/6B_COMB.seq:*/cgn2_6/ptodata/2/ina/6B_COMB.seq:* /cgn2_6/ptodata/2/ina/5A_COMB.seq: Issued_Patents_NA:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		Description	: .	seducince of Applit	Sequence 3, Appli	1,	7	, i	Sequence 3, Appli	1	11	11	11	1, 1	11,	Sequence 7, Appli	11	Ξ	Sequence 3, Appli	15,	15,	15,	Sequence 15, Appl	15,	15,	Sequence 10, Appl	20,	20,	20,	20.
		ID	770 - 00 - 446 - 000 - 0	02-00-440-00-00	PCT-US93-10034-3	US-07-940-605A-1	US-08-184-422-7	US-08-360-923A-1	US-08-431-055-3	US-08-690-096-1	US-08-249-189-11	US-08-484-624A-11	US-08-477-733B-11	US-08-763-995-1	US-09-088-913A-11	US-08-589-771B-7	US-08-769-819-11	US-08-770-974-11	US-08-858-197-3	US-08-249-189-15	-08-484-	US-08-477-733B-15	-09	-08-769	-770	-446-922	US-08-249-189-20	US-08-484-624A-20	-08-477	US-09-088-913A-20
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		Score	7.75	*	776.4	776.4	776.4	776.4	776.4	776.4	776.4	776.4	776.4	776.4	776.4	776.4	776.4	776.4	776.4	638.2	638.2	638.2	638.2	638.2	638.2	637.2	637.2		637.2	637.2
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Appl Appl	Appli Appli	Appl1 Appli	Appli	Appii Appli	Appli	App11	App11	Appl	Appl	Appl	Appl	Appl	Appl
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28	31	33	34	36	37.	38	39	40	4.1	42	43	44	45

ALIGNMENTS

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APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
TITLE OF INVENTION: Proteins
CORRESPONDENCE: 11.
                                                                                                                                                                                                                                                                                                                               COMPOTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: MICROSOft Word for Apple, Version 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446,922.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DAIL.
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/107,353
APPLICATION NUMBER: 08-13-93
                                                                                                                                                                                                                    E: Immunex Corporation 51 University Street
                                     Sequence 3, Application US/08446922 Patent No. 5716805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Perkins, Patricia A
RECIGSTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: CDNA to mRNA HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 786 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
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                                                                            GENERAL INFORMATION:
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STREET: 51.
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RESULT
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Proteins

FITLE OF INVENTION:

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CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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                                                                                                                                                             121 CITTITGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
                                                                                                                                                                                                           TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
                                                                                                                Sequence 3, Application PC/TUS9310034
GENERAL INFORMATION:
APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
                                                                              ;
0
                                                     Length 786;
                                                                              Indels
                                                                              9
                                                       Score 776.4; DB Pred. No. 2e-220;
                                                                               0; Mismatches
                                                       tch 98.8%;
al Similarity 99.2%;
780; Conservative
 CDS
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PCT-US93-10034-3
; NAME/KEY:
; LOCATION:
US-08-446-922-3
                                                        Query Match
Best Local S
Matches 780
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AACAAAGAGGAGACGAAGAAAAAAAAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 ATGATCGAAACATACAACCAAACTTCTCCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
                                                                                                                               COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  98.8%; Score 776.4; 99.2%; Pred. No. 2e-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No. 2e-2
0; Mismatches
                                                                                                                                                                                    APPLICATION NUMBER: PCT/US93/10034
FILING DATE:
                         Immuhex Corporation
                                     51 University Street
                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                         NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,69
                                                                                                                                                                                                                                                                                           TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                       OMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                  CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                     LENGTH: 786 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 99.2
Matches 780; Conservative
                                                                                                                                                                         CURRENT APPLICATION DATA:
NUMBER OF SEQUENCES: 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN: CD40-L
                                                     Seattle
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                                                                              USA
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                                                                                           98101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; LOCATION:
PCT-US93-10034-3
                            ADDRESSEE:
                                                                              COUNTRY:
                                                   CITY:
STATE:
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; LOCATION:
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361 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 420
                                                                                                                              CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                         GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG
                                                                           CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: ARUFFO, ALEJANDRO
APPLICANT: HOLLENBAUGH, DIANE
APPLICANT: LEDBETTER, JEFFRRY A.
TITLE OF INVENTION: SOLUBLE LIGANDS FOR CD40
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   E: Pennie & Edmonds
1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       UMBER: US/07/940,605A
04-SEP-1992
                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/07940605A Patent No. 5540926 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Misrock, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 562
TELECOMMUNICATION
TELEPHONE: 212 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 212 869-884/9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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TYPE: nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: New York COUNTRY: U.S.A.
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781 CTCTGA 786
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                                                                            1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
                                                                                                 22 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
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     Length 840;
                                         Indels
Score 776.4; DB 1;
Pred. No. 2.1e-220;
); Mismatches 6;
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99.2%;
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DAVISON, BARRY
                                           Conservative
                    Similarity
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APPLICANT: ARMITA
APPLICANT: DAVISO
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   Query Match
Best Local Simi
Matches 780;
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                                                      361 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAAACAACATCTGTGTTACAGTGG
                                                                                                                                                                                                                                                                   CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                  301 AACAAAGAGGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: Apple Macintosh System 7.1 SOFTWARE: Microsoft Word for Macintosh, Version #5.1a
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APPLICATION NUMBER: USSN 08/172,664
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51 UNIVERSITY STREET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: FANSLOW, MILLIAM
APPLICANT: LONGO, DAN L.
APPLICANT: MURPHY, WILLIAM
TITLE OF INVENTION: METHOD OF
TITLE OF INVENTION: DISEASE CI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: .. ARMITAGE, RICHARD
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CORRESPONDENCE ADDRESS:
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FILING DATE: Decemb
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| 826 CTCTGA 831
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                                                            DETECTION AND TREATMENT OF MUTATIONS
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Pred No. 2.1e-220;
0; Mismatches 6;
                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: Apple Operating System 7.1 SOFTWARE: MS WORD for Apple 5.1, Version a CORTENT APPLICATION DATA: APPLICATION NUMBER: US/08/184,422
                                                                                   IN A CD40 LIGAND GENE
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51 UNIVERSITY STREET
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APPLICATION NUMBER: 08/009,258
FILING DATE: 01/22/93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: PERKINS, PATRICIA ANNE REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 2816
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
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Similarity 99.2%;
0; Conservative
                        MELANIE
                                            WIDMER, MICHAEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                                       CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 800
                                          APPLICANT: WIDMER, P
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                       NUMBER OF SEQUENCES:
                                                                                                                                                                                                      WASHINGTON
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; LOCATION:
US-08-184-422-7
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Matches 78
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61 ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
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                                                                                             22 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACCTCCCCATCACC
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               CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                                                                                                                                                                                                                                                                                               APPLICANT: KEHRY, MERILYN R
APPLICANT: CASTLE, BRIAN E
TITLE OF INVENTION: MENDOS FOR PROLIFERATING AND
TITLE OF INVENTION: DIFFERENTIATING B CELLS, AND USES THEREOF
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Pred. No. 2.1e-220;
0; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: STERN, KESSLER, GOLDSTEIN & FOX
STREET: 100 NEW YORK AVE. N.W. SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: MILLMAN, ROBERT A
REGISTRATION NUMBER: 36,217
REFERENCE/DOCKET NUMBER: 1011.1030000/RAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/431,055
FILING DATE: 28-APR-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/234,580
FILING DATE: 28-APR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: PATENTIN Release #1.0, CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                Sequence 3, Application US/08431055 Patent No. 5817516
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99.2%;
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TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 3:
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Best Local Similarity 99.2
Matches 780; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: WASHINGTON
STATE: D.C.
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy d1
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; LOCATION:
US-08-431-055-3
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Pred. No. 2.1e-220,
0; Mismatches 6;
            NAME: Perkins, Patricia A.
REGISTRATION UNUBER: 34,693
REFERENCE/DOCKET UNUBER: 2818
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 840-base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 98.8%;
Best Local Similarity 99.2%;
Matches 780; Conservative (
ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
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US-08-360-923A-1
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                                                                 202 GAAGATTTGTATTGATGAAAAGGATACAGAGATGCAACACAGGAGAAAAGATCCTTATCC 261
                                                                                                 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
                                                                                                                481 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                                 361 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
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                                                                                                                                                              Version #1.25
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              equencolatera (Control of September of Seneral INFORMATION: APPLICANT: ARUFFO, ALEJANDRO APPLICANT: HOLLENBETTER, JEFRERY A. COLUGANT: LEDBETTER, JEFRERY A. COLUGANT: """PWTION: SOLUBLE LIGANDS F
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Patentin Release #1.0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/08690096 patent No. 5945513
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1155 Avenue of
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10036-2711
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802 CTCTGA 807
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JABER: US/08/690,096
31-JUL-1996
                                                             07/940,605
                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: 212 790-9090
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
                                                                                                        NAME: MISTOCK, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 56
                                                                                                                                                                                                                                                                                               DNA (genomic)
                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07
FILING DATE: 04-SEP-1992
                                                                               FILING DATE: 04-SEP-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                      LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                Matches 780; Conservative
                                                                                                                                                                                                                                                                                                                            CDS 22..807
                                                                                                                                                                                                                                                                                  linear
                                                                                                                                                                                                                                                                                                                                                                                                                  Similarity
                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                 TOPOLOGY: lin
                                                                                                                                                                                                                                                                                                                            NAME/KEY:
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US-08-690-096-1
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IMMEDIATE SOURCE CLONE: CD40-L
                                               ; NAME/KEY:
; LOCATION:
US-08-249-189-11
                                                                                                                                               Best Local Sin
Watches 780;
                  CLONE:
FEATURE:
                                                                                                                             Query Match
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                                                                                            622 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 681
                               661 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                              OPERATING SYSTEM: Apple Operating System SOFTWARE: Microsoft Word for Apple, versicurrent Application Data:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/249,189
                                                                                                                                                                                                                                                                                                                                                          APPLICANT: SRINIYASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 514

TOR APPLICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
CITY: SEATTLE
STATE: WASHINGTON
                                                                                                                                                                                                                                                                   Sequence 11, Application US/08249189
Patent No. 5961974
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: 07/783,70 FILING DATE: October 25, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Perkins, Patricia A. RECISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 2E TELECOMMUNICATION INFORMATION: 2065870430
                                                                                                                                                                                                                                                                                                                 APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                              APPLICANT: FANSLOW, WILLIAM APPLICANT: SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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ORIGINAL SOURCE:
                                                                                                                                                         781 CTCTGA 786
                                                                                                                                                                           111111
802 CTCTGA 807
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US-08-249-189-11
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                                                                                                                                                                                                                                                                                    226 GAAGATTTTGTATTCATGAAAGGATACAGAGATGCAACACAGGAGAAGATCCTTATCC 285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 661 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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                                                                                                                                                                    61 ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
                                             Gaps
                                                                                    1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
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                                                                                                                                                                                                                                                                                                                                                                                                                                          241 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTA
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  Length 840;
                                           Indels
.4; DB 2;
2.1e-220;
                      Pred. No. 2.16
0; Mismatches
Score 776.4;
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US-08-484-624A-11
'Sequence 11, Application US/08484624A
'Patent No. 5962406
98.8%;
                                           Conservative
                         Similarity
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                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: Apple Operating System 7.5.5 SOFTWARE: MS WOOT for Apple Power Macintosh, version 6.0.1 CURRENT APPLICATION DATA:
                                    APPLICANT: SRINIVASAN, SÜBHASHINI
APPLICANT: GIBSON, MARYLOU
APPLICANT: MORRIS, ARVIA E.
APPLICANT: MORRIS, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
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Pred. No. 2.1e-220;
0; Mismatches 6;
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APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
CLASSIFICATION:
                                                                                                                                                                       5: IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 08/477,
FILING DATE: June 07, 1995
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ilarity 99.2%;
Conservative
ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk COMPUTER: Apple Macintosh
                                PRIGGS, MELANIE
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IMMEDIATE SOURCE:
CLONE: CD40-L
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; LOCATION: 46..831
US-08-484-624A-11
                                                                                                                                                                                                                                WASHINGTON
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Best Local Similarity
Matches 780; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
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                                                                                                                                                                                                 STATE: WASTE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL:
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CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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61 AIGAAAAITITITAIGIATTTACITACIGITITICCITATCACCCAAAIGATIGGAICIGIG
                                                            121 CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT
                                                                                                                  GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC
                                                                                                                                                                          NOVEL CYTOKINE THAT BINDS CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FANSLOW, WILLIAM
SPRIGGS, MELANIE
SRINIVASAN, SUBHASHINI
GIBSON, WARYLOU
MORRIS, ARVIA E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11, Application US/08477733B
Patent No. 5981724
GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MCGREW, JEFFERY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: MCGREW, J
TITLE OF INVENTION: '
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          781 CTCTGA 786
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US-08-477-733B-11
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APPLICANT:
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406 CAAATTGCGCCACATGTCATAAGTGAGGCCAGCAGCAGTAAAACAACATCTGTGTTACAGTGG 465
                                                                                                                                                                                                                                                                                                                  481 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT 540
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
226 GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC 285
                                                                                   TTACTGAACTGTGAGGAGATTAAAAGCCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 345
                                                                                                                                                                                                                                                                                            GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 765
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                                                                                                                                                                                                           361 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
                                                                                                                            AACAAAGAGGAGAGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                           APPLICANT: MARASKOVSKY, EUGENE
TITLE OF INVENTION: METHOD OF ACTIVATING DENDRITIC CELLS
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SYSTEM: Apple Operating System 7.5.5 Microsoft Word for Macintosh, Version #6.0.1
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COUNTRY: USA
ZIP: 9101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Power Macintosh 7200/90
""" Apple Operating Symmetry: Apple Operating Symmetry: Apple Operating Symmetry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: USSN 08/677,762
FILING DATE: 10 JUL 1996
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APPLICATION NUMBER: US/08/763,995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
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CITY: SE
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MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/477,733B
FILING DATE: June 07, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 840;
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Pred. No. 2.1e-220;
0; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
FILING DATE: December 5, 1991
CLASSIFICATION 435
PRIOR APPLICATION AMBER: 07/83,707
                                                                                                                                                                                                                                             CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/249,189
FILING DATE: May 24, 1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: OCTOBER 25, 1991
CLASSIFICATION: 435
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APPLICATION NUMBER: 07/969,703
                                                                                                                                                                                                                                                                                                                                                                                                                                                           07/805,723
                                                                                                                                                                                                                                                                                                                                                                                             October 23, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34.628
REFERENCE/DOCKET NUMBER: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98.8%;
99.2%;
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TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 98.8
Best Local Similarity 99.2
Matches 780; Conservative
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  SEATTLE
WASHINGTON
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US-08-477-733B-11
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                                          USA
                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
                                                           98101
                                          COUNTRY:
                   STATE:
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GIGACIGAICCAAGCCAAGTGAGCCAIGGCACTGGCTICACGICCTIIGGCTIACICAAA 780
                                                                         SYSTEM: Apple Operating System 7.5.5
MS Word for Apple Power Macintosh, version 6.0.1
                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: MCGREW, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Operating System 7.5.5
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707.
FILING DATE: October 25, 1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                          Sequence 11, Application US/09088913A Patent No. 6087329
                                                                                                                                                                                                                                                                                                                                                                     SRINIVASAN, SUBHASHINI
GIBSON, MARYLOU
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 08/477, FILLME DATE: June 07, 1995 PRIOR APPLICATION DATA: APPLICATION NUMBER: 07/969,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                   RICHARD
                                                                                                                                                                                                                                                                                                                                    FANSLOW, WILLIAM
SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple O
                                                                                                                                                                                                                                                                                                                                                                                                          MORRIS, ARVIA E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: MS WORD for P
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
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TYPE: nucleic acid
STRANDEDNESS: single
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SEQUENCE CHARACTERISTICS:
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826 CTCTGA 831
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 776.4; DB 3;
Pred. No. 2.1e-220;
0; Mismatches 6;
                         2845-A
   REGISTRATION NUMBER: 34,693
REFERENCE/POCKET UNBER: 2845
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)2364
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 99.2%;
Matches 780; Conservative
                                                                                                                                      : 840 base pairs
nucleic acid
                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
                                                                                                                                                                             single
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46..831
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                                                                                                                                                                                                            MOLECULE TYPE:
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                                                                                                                                        LENGTH:
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                                                                                                                                                                                                                 Indels
                                                                                                                                                                      Score 776.4; DB 3;
Pred. No. 2.1e-220;
0; Mismatches 6;
                                                                                                                                                                      Query Match 98.8%;
Best Local Similarity 99.2%;
Matches 780; Conservative
  Homo sapiens
                                                                         ; NAME/KEY: CDS
; LOCATION: 46..831
US-09-088-913A-11
                IMMEDIATE SOURCE
CLONE: CD40-L
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ORGANISM:
                                                      FEATURE:
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                                                                          TREATMENT OF INDIVIDUALS EXHIBITING DEFECTIVE CD40L (as amended)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 776.4; DB 3;
Pred. No. 2.1e-220;
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99.2%; Pred. No. 2.1e
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                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: MS-DOS/Windows 95
SOFTWARE: Word for Windows 95, 7.0a
                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/589,771B
                                                                                                                                                : IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/009,258
                                                                                                                                                                                                                                                                                                                                                                                              January 22, 1996
                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: HENRY, JANIS C.
REGISTRATION NUMBER: 34,347
REFERENCE/DOCKET NUMBER: 25
TELECOMMUNICATION INFORMATION:
             RENSHAW, BLAIR
SPRIGGS, MELANIE
WIDMER, MICHAEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 99.2
Matches 780; Conservative
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INFORMATION FOR SEQ. ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 01/22/93
CLASSIFICATION: 435
                                                                                                                                CORRESPONDENCE ADDRESS:
                                                                  TITLE OF INVENTION:
TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              nucleic acid
                                                                                                                                                                                       COUNTRY: HC*
: FANSLOW,
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ORIGINAL SOURCE:
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MOLECULE TYPE:
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                                                                                                                                                ADDRESSEE:
STREET: 51
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                 APPLICANT:
APPLICANT:
                                                    APPLICANT:
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FEATURE:
APPLICANT
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; Sequence 7, Application US/08589771B

RESULT 13 US-08-589-771B-7

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541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 600
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                                                    241 TIACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
                                                                  CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 420
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GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACAACAGGAGAAAGATCCTTATCC 240
                 421 GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG
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APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT
NUMBER OF SEQUENCES: 24
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Microsoft Word for Apple,
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51 UNIVERSITY STREET
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Patent No. 6264951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
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OPERATING SYSTEM: Apple O
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MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION
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FILING DATE: October 23, 1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
ATTORNEY/AGENT INFORMATION:
                                                        APPLICATION NUMBER: 08/249,189
FILING DATE: May 24, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/484,624
FILING DATE: 07-JUN-1995
                                                                                                                                                                                                                                                                                               NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
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CLONE: CD40-L
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US-08-769-819-11
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                       Gaps
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                                           · 1 ATGATCGAAACATACAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
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Length 840;
  Score 776.4; DB 4;
Pred. No. 2.1e-220;
                        0; Mismatches
   98.8%;
ilarity 99.2%;
Conservative
               Similarity
                           780;
   Query Match
Best Local S:
Matches 780,
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CLONE:
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CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
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SOFTWARE: Microsoft Word for Apple, version 5.la
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APPLICANT: SPRIGGS, MELEANIE
APPLICANT: STRNYVASAN SUBHASHINI
APPLICANT: GIBSON, MARYLOU
APPLICANT: GIBSON, MARYLOU
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
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APPLICATION NUMBER: 08/477,733
FILING DATE: 02-AUG-1995
APPLICATION NUMBER: 08/249,189
FILING DATE: MAY 24, 1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3: IMMUNEX CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 07/783,70 FILING DATE: October 25, 1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
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CITY: SEATTLE
STATE: WASHINGTON
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
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661 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 776.4; DB 4;
Pred. No. 2.1e-220;
0; Mismatches 6;
TELECOMMUNICATION INFORMATION
TELEPHONE: 2065870430
TELEPAX: 2065870606
INPORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                98.8%;
ilarity 99.2%;
Conservative
                                                                                                                                            STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: HOMO SapienS
IMMEDIATE SOURCE:
                                                                                                       LENGTH: 840 base pairs
TYPE: nucleic acid
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⁷⁸¹ CTCTGA 786 | | | | | | | | 826 CTCTGA 831 Oy Op

Search completed: May 30, 2002, 02:51:32 Job time: 11596 sec

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Result
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Perfect score:
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				iers	Qualif	Location/Qualifiers	ĭ		FEATURES
			•	Patent: US 5817516-A 1 06-OCT-1998;	-A 1 0	5817516	: US	Patent	JOURNAL
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				• =		ed.	sifie	Unclassified	
						٠		Unknown	ORGANISM
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							78	AR044778	ACCESSION
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Best Matches

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1 (bases 1 to 1250)
Tripp, R.A., Anderson, L.J. and Brown, M.P.
Cd40 ligand adjuvant for respiratory syncytial
Patent: WO 0156602-A 1 09-AUG-2001;
The Secretary, Department of Health and Human S
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Eukaryota;
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                            Armitage, R.J., Fanslow, W.C., Spriggs, Gibson, M.G., Morris, A.E. and McGrew, J DNA encoding CD40 ligand, a cytokine Patent: US 5981724-A 1.09-NOV-1999;
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Armitage, R.J., Fanslow, W.C. and Spriggs, M. CD40 ligand polypeptide
Patent: US 6087329-A 1 11-JUL-2000;
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Submitted (07-APR-1992) M.K.
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Mammalia; Eutheria;
1 (bases 1 to 1250)
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                                                                                                                                                                                                                                                                                                                                                                                                                                       /product="CD40 surface protein"
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/translation="MIETYSQPSPRSVATGLPASMKIFMYLLTVFLITQMIGSVLFAV
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/codon_start=1
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/db_xref="taxon:10090"
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                                                                                                                                         Direct Submission
Direct (14-JUL-1997) Immunology, King's Co
Submitted (14-JUL-1997) Immunology, King's Co
                                                                                                                                                                                            2 (bases 1 to 783)
Daniel, K.C., Foss, Y., Mous.
Farzaneh, F. and Gaken, J.A.
                                                                                                                                                                                                                                                    Daniel, K.C., Foss, Y., Moussavi, A., Macary Farzaneh, F. and Gaken, J.A. Cloning and sequencing of rat CD40 ligand
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                             /tissue_type="spleen"
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                                                          /organism="Rattus norvegicus"
/strain="PVG"
/db_xref="taxon:10116"
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               codon_start=
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                                                                                               TCCATTCACTTGGGCGGAGTATTTGAATTACAAGCAGGTGCTTCTGTGTTTGTCAACGTG
                                                                                                              TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTCTGTGTTTTGTCAACGTG
   TGA 783
                               ACTGAAGCAAGCCAAGTGATCCACGGAATTGGCTTCTCATCTTTTGGCTTACTCAAACTC
                                                              ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC
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/db_xref="GI:4102614"
/translation="MIETYSQPSPRSVATGLPASMKIFMYLLTVFLITOMIGSVLFAV
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/thrresplsorffivslmlkpssgserillraanthsssklce
/qosihlgvfelqagasyfvnnyeasgvihgigfssfgllkl"
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                                                                                                                                                                                                                                                                                             GAAGATTTTGTATTCATAAAAAAACCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
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                                                                                                                                                                                               CTTTTTGCTGTGTATACTAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180
                                                                                                                                                                                                                                               ATGAAGATTTTTATGTATTTACTTACTGTTTTTTCTTATCACCCAGATGATTGGGTCGGTG
AACAAAGAAGAAGAAAAAAAAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAA
                                                                                                           TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGÀTATAACGTTA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Rattus norvegicus CD40 ligand
AF116582
AF116582.1 GI:4545249
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Hallett, K.M. and Oa
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2900 W. Oklahoma Ave., Milv
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                           /product="CD40 ligand"
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TVKREGLLYVVTQVTGVTGCSNREPLSQRPFIVSLMLKPSSGSERILLRAANTHSSSKLCE
QOSIHLGGVFELQAGASVFVNVTEASQVIHGIGFSSIGLLKL"
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Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /strain="Sprague Dawley"
/db_xref="taxon:10116"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /organism="Rattus norvegicus"
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l mRNA, complete cds.
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 AGTAATGCAGCATCCGTTCTACAGTGGGCCAAGAAAGGATATTATACCATGAAAAGCAAC 450
                      GAAATGCAAAGAGGTGATGAGGATCCTCAAATTGCAGCACACGTTGTAAGCGAAGCCAAC
                                 GAAATGCAAAGAGGTGATGAGGATCCTCAAATTGCAGCACACGTTGTAAGCGAAGCCAAC 390
                                                                                                          AGATGCAACAAAGGAGGAGCTTTATCCTTGCTGAACTGTGAGGAGGATGAGAAGGCAA
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                                                                                                                                                                                              632;
                                                                                                                                                                                                                                                                                                                                                                                               Sequence 22
AR076933
AR076933.1
                                                                                                                                                                                                                                                                                                   Armitage, R.J., Fanslow, W.C. and Spriggs, M.K. Monoclonal antibodies to CD40 ligand, pharmaceutical composition comprising the same and hybridomas producing the same patent: US 5961974-A 22 05-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                   Unknown.
                                                                                                                                                                                                                                                                                                                                                                                                                               AR076933
                                                                                                                                                                                                                                                                                                                                                       Unclassified
                                                                                                                                                                                                                                                                                                                                                                            Unknown.
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178 c 203 g
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patent US 59
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Pred. No. 2.6e-153;
0; Mismatches 1;
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                                                                                                                                                                                            Gaps
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RESULT 14
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ACCESSION
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AR078323
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AR085426
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Search completed: May 30, 2002, 02:48:59 Job time: 22018 sec

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                      AL232112 Tetraodon
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ALIGNMENTS

DEFINITION ACCESSION VERSION KEYWORDS REFERENCE AUTHORS RESULT BF599437 FEATURES COMMENT SOURCE Locus TITLE ORGANISM MEDLINE JOURNAL source Contact: Smith TPL
USDA, ARS, US Meat Animal Research Center
PO Box 166, Clay Center, NE 68933-0166, USA
Tel: 402 762 4366
Fax: 402 762 4390
Email: smith@email.marc.usda.gov
Single pass sequencing. Bases called and all
v0.980904.e. Vector identified by cross_matv (bases 1 to 492)
Smith, T.P.L., Grosse, W.M., Freking, B.A., Roberts, A.J., Stone, R.T., Casas, E., Wray, J.E., White, J., Cho, J., Fahrenkrug, S.C., Bennett, G.L., Heaton, M.P., Laegreid, W.W., Rohrer, G.A., Chitko-McKown, C.G., Pertea, G., Holt, I., Karamycheva, S., Liang, F., Quackenbush, J. and EST. BF599437 BF599437.1 GI:11695919 Sequence evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle Genome Res. 11 (4), 626-630 (2001) BF599437 492 bp mRNA linear 263218 MARC 3BOV Bos taurus cDNA 5', mRNA sequence Bos taurus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea; Seq primer: ATTTAGGTGACACTATAG. BACKWARD: GTTTTCCCAGTCACGACG Plate: 33 row: N column: 5 FORWARD: AGGAAACAGCTATGACCAT PCR PRimers 21180013 Keele, J.W. Bovidae; Bovinae; Bos. -minmatch 12 options. /organism="Bos taurus' /db_xref="taxon:9913" Location/Qualifiers called and alt_trimmed with phred cross_match with the EST 25-APR-2001 -minscore

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  365;
                                                                                                                   Smith, T.P.L., Grosse, W.M., Freking, B.A., Roberts, A.J., Stone, R.T., Casas, E., Wray, J.E., White, J., Cho, J., Fahrenkrug, S.C., Bennett Casas, E., Watton, M.P., Laggraid, W.W., Rohrer, G.A., Chitko-McKown, C.G., G.L., Haatton, M.P., Laggraid, W.W., Rohrer, G.A., Chitko-McKown, C.G., Pertea, G., Holt, I., Karamycheva, S., Liang, F., Quackenbush, J. and
                                                                                                                                                                                                                                                                        EST
                                                          Sequence evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle Genome Res. 11 (4), 626-630 (2001)
                                                                                                                                                                                                                                            Bos taurus
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75217 MARC
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                                                                                                                                                                                                                                                                                        AW486605.1
                              Contact: Smith TPL
                                              21180013
                                                                                                       Keele,J.W.
                                                                                                                                                                                                            Mammalia;
DA, ARS, U
Box√166,
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103 c 104 g 122 t
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/tissue_type="pooled"
/lab_host="DH10B"
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to
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 US Mea
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Eutheria; Cetartiodactyla; Ruminantia; Pecora;
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Pred. No. 2.7e-54;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: smith@email.marc.usda.gov single pass sequencing. Bases called and trimmed with phred v0.980904.e. Vector identified by cross_match with the -min:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BACKWARD: GTTTTCCCAGTCACGACG
Plate: 34 row: F column: 23
Seq primer: ATTTAGGTGACACTATAG
                                                                                                                                                                                                                           AI982044 638 bp mRNA linear EST 07-M pat.pk0072.c9.f chicken activated T cell cDNA Gallus gallus clone pat.pk0072.c9.f 5' similar to CD40 ligand, mRNA sequen AI982044
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                                                                                                               FORWARD: AGGAAACAGCTATGACCAT
             Tirunagaru, V.G., Sofer, L., Cui, J. and Burnside, J. an expressed sequence tag database of T-cell-enriched chicken splenocytes: sequence analysis of 5251 clones Genomics 66 (2), 144-151 (2000)
                                                                                                    Archosauria; Aves; Ne Phasianinae; Gallus.
                                                                                                                                                                                              EST
                                                                                                                                                                             chicken
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402 762 4390
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17 c 82 g 105 t
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/clone_lib="MARC 1BOV"
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/lab_host="DH10B"
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Pred. No. 3.1e-42;
0; Mismatches 51
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fly), genomic survey sequence.
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                                  Submitted (02-JUN-1999) Genoscope - Centre National de Sequencage BP 191 191006 EVRY cedex - FRANCE (E-mail: seqref@genoscope.cns.fr - Web: www.genoscope.cns.fr)
Determination of this BAC-end sequence was carried out as part of collaboration with the Berkeley Drosophila Genome Project (BDGP). The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further information
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CNS005TE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Email: joan@UDel.Edu, www.chickest.udel.edu
Seq primer: T7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Joan Burnside Molecular Endocrinology
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40 Townsend Hall, Newa
                                                                                                                                                                                      Direct Submission
                                                                                                                                                                                                           Genoscope.
                                                                                                                                                                                                                                              Eukaryota: Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilldae; Drosophila.
melanogaster
                   please see http://www.fruitfly.org The BDGP Drosophila
                                                                                                                                                                                                                                                                                                                                       ruit fly.
                                                                                                                                                                                                                             (bases 1 to 997)
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/lab_host="E.coli TOP10 F'"
/note="Vector: pcDNA3"
171 c 151 g 180 t 4 others
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/clone_lib="chicken activated T cell cDNA"
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library was prepared by Kazutoyo Osoegawa and
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Pred. No. 2.2e-05;
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                                                                                                                                            segref@genoscope.cns.fr
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GAGARRGRRRRGRARRAARGAARRRARRARRRRRRAAARRGAGRAGARARGR
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                                                                                                                                                                   GAGAARRRGRRGRGAGARRRGRGRRRRGAGARRRRRGRRMRTRRARRRRRRRAGAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       and how to order individual BAC clones, the entire library, or filters for hybridization from the BACPAC Resource Center can be found at http://bacpac.med.buffalo.edu/drosophila_bac.htm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPCI-98 and was constructed by partial EcoRI digestion of Drosophila DNA provided by the BDGP from the Liogenic Strain y2; cn bw sp, the same strain used for the BDGP's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         P1 and EST libraries.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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/db_xref="taxon:7227"
/clone_lib="RPCI-98"
/clone="BACR12K22"
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                                                                                                                                                  AUTHORS
TITLE
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                                                                                                                               Li,W.B., Gruber,C., Jessee,J. and Polayes,D. Full-length cDNA libraries and normalization Unpublished (2001)
                                                                                                                                                                                                                                                                         EST.
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AL534423 LTI_FL013_FBrn1
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AL534423
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                                                                                                 Genoscope - Centre National de Sequencage
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                                                                segref@genoscope.cns.fr,
                                                                                91006 EVRY cedex - France
/organism="Homo sapiens"
/db_xref="taxon:9606"
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AUTHORS
TITLE
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Maryland 20850, USA Fax: (1) 301 610 8371 Email:
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/dev_stage="pooled_tissue fro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     fliang@lifetech.com URL :
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38.9%;
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10 week and 26 week)"
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Determination of this BAC-end sequence
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Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
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    Web: www.genoscope.cns.fr)
    Determination of this BAC-end sequence was carried out as part of a

                             Submitted (23-JUL-1999) Genoscope - Centre BP 191 91006 EVRY cedex - FRANCE (E-mail :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAGAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAAATTGCAGCACACGTTG 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAAAAWWMWCTTTGCMCTGAWTRTWGTTRTWTWAWWWMRTTRAAWWWMAAWWWMAAWWW
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please see http://www.fruitfly.org The BDGP Drosophila melanogaster BAC library was prepared by Kazutoyo Osoegawa and Aaron Mammoser in Pieter de Jong's laboratory in the Department of Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY The library is named RPCI-98 and was constructed by partial EcoRI digestion of Drosophila DNA provided by the BDGP from the isogenic strain y2; cn bw sp, the same strain used for the BDGP's Pl and EST libraries. A more detailed description of the library
                                                                                                                                                                                         Web: www.genoscope.cns.fr)

Determination of this BAC-end sequence was carried out as collaboration with the Berkeley Drosophila Genome Project The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further informat melanogaster genome using these BACs. For further informat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fly), ge
AL063921
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                                                                                                                                                                                                                                                                                                                                   Submitted (02-JUN-1999) Genoscope - Centre BP 191 91006 EVRY cedex - FRANCE (E-mail :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.

1 (bases 1 to 1101)
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/clone="BACN04E04"
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/plasmid="pBeloBAC11"
/db_xref="taxon:7227"
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segref@genoscope.cns.fr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             122 TTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCATG 181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       664 WTTTDTDTDDWDKRDRRR 647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DADDDTDGTKDDDDKDKWDDWDKAKGTWGDATWAWAATDWWWWGWADADWWTWDAAADDW
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Entamoeba histolytica
                                                                                                                                                                                                                                                                                                                                                    genomic, DNA sequence
                                                                                         Department of Eukaryotic Genomics
                                                                                                                  Contact: Brendan J Loftus
                                                                                                                                                     Determination of clone end sequences HM1:IMSS sheared DNA library
                                                                                                                                                                                                                                  Eukaryota; Entamoebidae;
                                                                                                                                                                                                                                                                                                               AZ691914.1 GI:11829180
                                                                                                                                                                                            Loftus, B., Van Aken, S. and Fraser, C
Email:, bjloftus@tigr.org
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                                                            Institute for Genomic Research
2 Medical Center Dr , Rockville,
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                                                          Center Dr.,
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Matches 137; Conserv
                                                                   AUTHORS
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                                                                                                                                                                                                                                                                                                                                                          308. AAGAGAAAAAAGAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       188 TTGTATTCATAAAAAAGCTAAAGAGATGCAACAAAGGAGGAACGATCTTTATCCTTGCTGA 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       68 TTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTGCTTTTTG 127
                                                                                                                                                                                                                                                                                                                                      ACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAG 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        High quality sequence start: High quality sequence stop: 3
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                                                                                                                                                                                                  423, bp DNA 11
"IM0229106F Mouse 10kb plasmid UUGC1M library clone UUGC1M0229106 F, DNA sequence.
A2438784
                                                                                                                                       Mus musculus
                                                                                                                                                                                      AZ438784.1
Mouse whole genome scaffolding with paired end reads from 10kb
                                                                Dunn, D., Aoyagi, A.,
                                                                                                                     Eukaryota; Metazoa;
                                                                                                                                                       house mouse
                 and Wright, D., Weiss, R.
                                                     Islam,H.,
                                                                                                   Mammalia; Eutheria;
                                                                                     (bases 1 to 423)
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                                Rose, M:, Rose, R., Stokes, R:,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          77.450.). The DNA was mechanically sheared to give a tight size distribution (~2 kb). The v + i method used for the library construction is described in detail in Smith.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            method for isolate identification.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          whole genome shotgun sequencing | sequencing: A Practical Approach,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Institute for Genomic Research (TIGR), Rockville,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /clone_lib="Entamoeba histolytica Sheared DNA"
/note="Vector: pHOS1; Site_1: Bst I; Construct
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /organism="Entamoeba histolytica"
/strain="HM1:IMSS"
/db_xref="taxon:5759"
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                                                 Longacre, S., Mahmoud, M.,
                                                                                                                                                                                         GI:10562797
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c 131 g 381 t
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                                                                  Barber, M.,
                                                                                                           Rodentia;
                                                                                                                           Chordata;
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                                                                                                                         Craniata; Vertebrata; Euteleostomi;
                                                                                                         Sciurognathi; Muridae;
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                                                                          Beacorn, T.,
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Press, 1999)."
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                                                         Meenen, E.,
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sculus genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
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                                                                                                                                                                                                                                 ATAGGAGAGTTGGGAAGGAGAAAAGAGAGAGAAAATGGTGTGATTATATATAAT
                                                                                                                                                                                                                                                                        AAAGGAGAAGGATCTTTATCCTTGCTGAACTGTGAGGAGGATGAGAAGGCAATTTGAAGAC 279
                                                                                                                                                                                                                                                                                                               GACTATGTAACAATAATTAATAAAAAAGAGTCCATGAATGTGAAAGATAGCAAGAAGGGGGT
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prime, m
AL514085
              AL514085 LTI_NFL006_PL2 Homo sapiens cDNA clone CL0BA004ZE10 prime, mRNA sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tel: 801 585 5606
Fax: 801 585 7177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Robert B. Weiss
University of Utah Genome Center
University of Utah
Rm. 308, Biomedical Polymers Research Bldg.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unpublished (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Email: ddunn@genetics.utah.edu
Insert Length: 10000 Std Err
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lass: plasmid ends
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      adaptored DNA was purified and size-selected for a 9.5 to 10.5 kb range using preparative agarose gel electrophoresis. Vector DNA was prepared from a derivative of pNPA2 (gil4732114|gb|AF129072.1), a copy-number inducible derivative of plasmid R1. The vector was ligated with adaptors complementary to the insert adaptors and purified. The sheard, adaptored mouse DNA was annealed to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          adaptored vector DNA, and transformed into chemically-competent E. coli XL10-Gold (Stratagene) cells and selected for ampicillin resistance."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               was blunt end-repaired with T4 DNA polymerase and T4 polynucleotide kinase. Adaptor oligonucleotides were ligated to the blunt ends in high molar excess. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (http://www.jax.org/resources/documents/dnares/). The DNA was hydrodynamically sheared by repeated passage through 0.005 inch orifice at constant velocity. The sheared DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     musculus C57BL/6J (male) was obtained from the Jackson
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/strain="C57BL/6J"
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TITLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note="Vector: pcMvSPORT 6; Site_1: NotI; 1st strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the pcMvSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies. Contact: Feng Liang Life Technologies, davision of Invitrogen 9800 Medical Center Drive Rockville, Maryland 20850, USA Fax: (1) 301 610 8371 Email: filang@lifetech.com URL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   http://fulllength.invitrogen.com"
13 c 3 g 207 t 4
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/clone_lib="LTI_NFL006_PL2"
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 46.8;
Pred. No. 6
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bp DNA linear GSS 15-MAY-2000
  survey sequence PUC-Ori end of clone
aodon nigroviridis, genomic survey
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GSS; genome survey sequence. Tetraodon nigroviridis. Tetraodon nigroviridis

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;

Actinopterygii;

Tetraodon nigroviridis genome surve 004012 of library G from Tetraodon

802 bp

sequence. AL232112

AL232112.1

GI:7891247

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AUTHORS
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                 ORGANISM
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTGCTTTTTG 127
                                                                                                                                                                                                                                                                                                                                                                                                   TIGTATICATAAAAAAGCTAAAGAGATGCAACAAAGGAGGAGGATCTTTATCCTIGCIGA 247
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCATGAAGATT 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAGAGAAAAAGAAAACAGCTTTG
                                                                                                                                                                                                                                                                                                                                   ACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAG 307
                                                                                                                                                                                                                                    AAAAAAAAGAAGAAGRAGTTKTR 100
                                                                                                                                                                                                                                                                                                   AGAGGTGTGTAGTGAGGGGGGGTGAATGTGTAGGGTRARATGTGRAGAAAARARAGGGGAA 124
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (12-APR-2000) to the EMBL/GenBank/DDBJ databases This sequence is a single read and was generated as part of a large scale clone-end sequencing project of the Tetraddon nigroviridis genome. For more information, please take a look at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Saurin, W. and Weissenbach, J.
Human gene number estimate provided by genome wide
Tetraodon nigroviridis DNA sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Unpublished
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                                                                                                                  Drosophila melanogaster genome sur BACN06A02 of DrosBAC library from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           http://www.genoscope.cns.fr/Tetraodon
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Direct Submission
                                                                               fly), genomic survey sequence.
AL100719
                                                                                                                                                CNS011TX
Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda;
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                                   fruit fly.
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/clone_lib="G"
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/db_xref="taxon:99883"
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Pred: No. 11;
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                                                                                                                  Drosophila melanogaster (fruit
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Bernot,A. and
                                                                                                                                      T7 end
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    254 AGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAGAAGAAGAGA 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     194 TCATAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCCTTGCTGAACTGTG 253
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Muscomorpha; Ephydroidea; Drosophilidae; Drosophila
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Submitted (23-JUL-1999) Genoscope - Centre National de Sequencage BP 191 91006 EVRY cedex - FRANCE (E-mail : seqref@genoscope.cns.fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Direct Submission
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Li,W.B., Gruber,C., Jessee,J. a Full-length cDNA libraries and Unpublished (2001)
                                                                                                                                                  prime, mRNA sequence.
AL514085
AL514085.1 GI:12777579
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AL514085 LTI_NFL006_PL2
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                                                  Eukaryota; Metazoa; Mammalia; Eutheria; 1 (bases 1 to 458)
                                                                                                                                    EST
                                                                                                      Homo sapiens
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/clone="BACN06A02"
/note="end : T7"
a 127 c 20 g
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/plasmid="pBeloBAC11"
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39.7%;
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                                                                                     Chordata; Craniata; Vertebrata;
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Pred. No. 11;
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                                                                          Hominidae;
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Best Local Similarity 41.1
Matches 113; Conservative
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402 ΑΑΑΑΑΑΑΑΤΑΤΑΑΑΑΑΑΑΛΛΛΛΛΛΛΛΛΛΛΛΛΛΛΛΛΑΑΑ 436
                       308 AAGAGAAAAAAGAAAACAGCTTTGAAATGCAAAGA 342
                                                                                188 TTGTATTCATAAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCCTTGCTGA 247
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Contact: Genoscope
Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - France
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note="Vector: pcMVSPORT 6; Site_1: Not1; 1st strand cDNA was primed with a Not1-oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the pcMVSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies. Contact: Feng Liang Life Technologies, a division of Invitrogen 9800 Medical Center Drive Rockville, Maryland 20850, USA Fax: (1) 301 610 8371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            http://fulllength.invitrogen.com"
13 c 3 g 207 t 4
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/db_xref="taxon:9606"
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lone_lib="LTI_NFL006_PL2"
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41.1%; Pred. No. 14;
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Search completed: May 30, 2002, 01:00:10 Job time: 22220 sec

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Thu May 30 05:46:11 2002
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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- nucleic search, using sw model OM nucleic Run on:

May 30, 2002, 03:03:15; Search time 680.87 Seconds (without alignments) 1974.450 Million cell updates/sec

US-08-982-272-5 783 Title: Perfect score: Sequence:

1 ATGATCGAAACATACAACCA.....TTGGCTTACTCAAACTCTGA 783

Scoring table:

IDENTITY_NUC Gapop 10.0 , Gapext 1.0

1736436 seqs, 858457221 residues Searched:

Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

N_Geneseq_032802:

/SIDS1/qcqdata/hold-geneseq/genesegn-embl/NA2002.DAT:* /SIDS1/gcgdata/hold-geneseq/geneseqn-emb1/NA2000.DAT:* /SIDS1/gcgdata/hold-geneseq/geneseqn-emb1/NA2001A.DAT: /SIDS1/gcgdata/hold-geneseq/geneseqn-emb1/NA2001B.DAT: /SIDS1/gcgdata/hold-geneseq/genesegn-embl/NA1996.DAT: /SIDS1/qcqdata/hold-geneseq/geneseqn-emb1/NA1999.DAT: /SIDS1/gcgdata/hold-geneseq/geneseqn-embl/NA1989 /SIDS1/gcgdata/hold-geneseq/genesegn-embl/NA1993 /SIDS1/gcgdata/hold-geneseq/genesegn-embl/NA1994 /SIDS1/gcgdata/hold-geneseq/genesegn-embl/NA1995 /SIDSI/gcgdata/hold-geneseq/geneseqn-embl/NN1983. /SIDSI/gcgdata/hold-geneseq/geneseqn-embl/NN1984. /SIDSI/gcgdata/hold-geneseq/geneseqn-embl/NN1985. /SIDSI/gcgdata/hold-geneseq/geneseqn-embl/NN1987. /SIDSI/gcgdata/hold-geneseq/geneseqn-embl/NN1987. /SIDSI/gcgdata/hold-geneseq/geneseqn-embl/NN1987. /SIDS1/gcgdata/hold-geneseq/genesegn-embl/NA199 /SIDS1/gcgdata/hold-geneseq/geneseqn-embl/NA199 /SIDS1/gcgdata/hold-geneseq/genesegn-embl /SIDS1/gcgdata/hold-geneseq/genesegn-emb /SIDS1/gcgdata/hold-geneseq/genes//SIDS1/gcgdata/hold-geneseq/genes /SIDS1/qcqdata/hold-qeneseq/ /SIDS1

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

+[, ,				
NO.	Score	Match	Match Length DB ID	DB	ID	Description
1	783	100.0		19		Exemplary CD40 lie
7	773.4			19	AAV38999	Exemplary CD40 lie
e	760.6				AAV38998	CD40 ligand gene
4	760.6				AAV61062	Murine CD40 ligano
S	759			15	AAQ63960	Mouse CD40-L type
9	759					CD40 ligand coding
7	759					Mouse CD40-L codir
œ	754.8		782	14	AAQ41507	Murine CD40-L DNA
σ	6.45					Mind and ide

Murine CD40 ligand Exemplary nucleoti	Mouse trimeric CD4	Exemplary CD40 lig	Human CD40-L type	CD40 ligand gene u			CD40 ligand gene.	Human CD40 ligand	cDNA of CD40L, a n	Human CD40 ligand	Human CD40-L codin	Nucleotide sequenc	Human CD40 ligand	DNA encoding novel	Human CD40L mutein	Genomic sequence o	Human CD40-L CDNA.	· Exemplary CD40 lig	D40	CD40 ligand gene u		CD154	CD154	CD154	CD154	CD154	CD154	Canine CD154 cDNA	ectide segu	-1 gp120 V3	-	٠,	gbI	Human CD40-L/Fc fu	
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ALIGNMENTS

AAV39001 standard; DNA; 783 BP. 23-SEP-1998 (first entry) AAV39001; RESULT AAV39001

Exemplary CD40 ligand gene used in the course of the invention.

CD40 ligand; alteration; immunoreactivity; human cell; accessory molecule ligand; AML; gene therapy; treatment; neoplasia; autoimmune disorder; rheumatoid arthritis; vaccine; chimera; ss.

Chimeric - Mus sp. Chimeric - Homo sapiens

WO9826061-A2.

18-JUN-1998

.97WO-US22740. 08-DEC-1997; 97US-0982272. 01-DEC-1997; 09-DEC-1996;

(REGC) UNIV CALIFORNIA.

Sharma S; Cantwell M, Kipps TJ,

WPI; 1998-348521/30

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The present sequence represents an exemplary CD40 ligand gene, comprising nucleotides encoding the extracellular domains (Domains III and transmembrane domain (Domain II) the murine CD40 ligand gene (AAV38997) operatively linked to nucleotides encoding the cytoplasmic channer of the human CD40 ligand gene (AAV38988). The sequence is used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells. Vectors containing the AML genes can be used in gene through arthritis. They can applied to the containing the AML genes can be used in gene through arthritis. They can applied the containing the amount of sorders such as rheumatoid arthritis. They can applied the containing the contai for vaccination to produce immunity against a virus cell; οŧ Vectors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatment neoplasia or autoimmune disorders, e.g. rheumatoid arthritis Sequence 783 BP; 243 A; 161 C; 174 G; 205 T; 0 other; bacteria, protein, fungus or neoplasia. Disclosure; Page 106; 167pp; English.

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Length 783;
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 100.0%; Score 783; DB 19;
100.0%; Pred. No. 3.1e-208;
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comprising nucleotides encoding the extracellular domains (Domains III comprising nucleotides encoding the extracellular domains (Domains III) and IV) of the murine CD40 ligand gene (AAV38997) operatively linked to nucleotides encoding the extracellular domains III and IV) and transmembrane domain (Domain III) of human CD40 ligand gene (AAV38998). The sequence is used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of specification describes a introducing a gene encoding an accessory human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells. Vectors containing the AML genes can be used in
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Pred. No. 1.5e-205;
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99.28;
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96US-0032145.
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Chimeric - Homo sapiens.
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Pred. No. 5.4e-202;
); Mismatches 14; I
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                                                                                ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC 780
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                                                                    TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
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                                                                                                                                                                                                                                                                                                                                                Murine; mouse; CD40 ligand; TNF receptor family; activated T cell; type 2 membrane glycoprotein; cell.proliferation; differentiation; B cell; ds.
 gagccttcgagtcaacgcccattcatcgtcggcctctggctgaagcccagcattggatct
                         GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
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                                                                                                                                                                                                                                                                                                                                 DNA sequence
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13..795
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                                                                                                                                                                                                                                                            AAV61062 standard; DNA; 818 BP.
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numbers, which is a vast increase compared
                                                 Length
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                         G; 213 T; 0 other;
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                                                 Score 760.6; DB 19
pred. No. 5.5e-202;
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98.2%;
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                                                     Query Match
Best Local Similarity 98.2%
Matches 769; Conservative
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   increase in cell culturing methods
                            Sequence 818 BP;
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                                                                                                                                                                              /note= "nucleotides 148-780 code for the extracellular region (amino acids 50-260)"
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                                             Leucine zipper; trimerisation; trimeric CD40-L; fusion protein. hetero-oligomer; homo-oligomer; type II transmembrane protein; soluble CD40-L; tumour necrosis factor family; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                            Prepn. of soluble oligomeric mammalian proteins – using host cells to express a fusion protein comprising a leucine zipper domain and a heterologous mammalian protein
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                      CD40-L type II transmembrane protein coding sequence.
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Pred: No. 1.5e-201;
0; Mismatches 15;
                                                                                                                                                                 mouse_CD40-L
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                                                                                                                            Location/Qualifiers
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13-AUG-1993;
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Leucine zipper; fusion protein production; soluble oligomeric protein; heterologous maimmalian type II transmembrane protein; activated T cell; heterologous mammalian type II transmembrane protein; antibody production; CD40-L; B-cell proliferation; CD27-L; lymphocyte antigen; ss.
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aacaaagaagaaaaaaagaaaacagctttgaaatgcaaagaggtgatgaggatcctcaa
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                                                                                                                                                                           AAV12853 standard; cDNA to mRNA; 783
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/*tag= a
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91US-0805723
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25-OCT-1991;
05-DEC-1991;
23-OCT-1992;
13-AUG-1993;
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us-08-982-272-5.rng

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This sequence is the coding sequence for the mouse CD40 ligand (CD40-L). The encoded protein can be used in a fusion protein produced using the mathod of the invention. The method is for preparing soluble oligomeric protein by culturing a host cell transfected with a vector for the fusion protein. The soluble oligomeric proteins comprise a leucine zipper fused to the N terminus of the extracellular region of a heterologous mammalian type II transmembrane protein or to the C terminus of the extracellular region of a heterologous mammalian type II transmembrane extracellular region of a heterologous mammalian type II transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                          stimulates B-cell proliferation and antibody production in
                                                                                                                                                                                                                                                                                                         protein, where the leucine ripper is a peptide comprising at least part of AAW41171 or AAW41772, optionally with conservative amino acid substitutions, provided that the peptide trimerises in solution. Assubble fusion protein comprising the leucine zipper of AAW41171 linked to the extracellular region of CD40-L (a type II transmembrane protein that is found on activated T cells and acts as a ligand for the B-cell.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 aacaaagaagaaaaaaaaaaaaaaaaaggtttgaaatgcaaagaggtgatgaggatcctcaa 360
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                                                                                                                                                                                                                                                                                                                                                                                                                      antigen CD40) stimulates B-cell proliferation and antibody production a similar manner to membrane-bound CD40-L. A soluble fusion protein comprising the leucine zipper of ABW41172 linked to the extracellular region of CD27-L (a type II transmembrane protein that binds to the region of CD27-L (a type II transmembrane protein that binds to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        leucine zipper fused
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           transmembrane protein that binds nibits binding of CD27c (a fusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         comprising the extracellular region of CD27 and a human 1961
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 783;
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Pred. No. 1.5e-201;
0; Mismatches 15;
                                                                                                                       to extracellular region of transmembrane protein
                                                                                                          comprising
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EBV-transformed B cells expressing CD27-L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ymphocyte antigen CD27) inhibits binding
                                                                                                                                                         Example 1; column 23-24; 21pp; English.
                                                                                                               Soluble oligomeric fusion proteins
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nilarity 98.1%;
Conservative 0
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                                  Srinivasan
(IMMV) IMMUNEX CORP.
                                                                WPI: 1998-144799/13.
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nes 768; Conserv
                                                                              P-PSDB; AAW41179
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Best Local S
Matches 768
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This sequence encodes the mouse CD40 receptor ligand (CD40-L). The hyportion relates to anti-human CD40-L monoclonal antibodies M90 secreted by hybridoma hCD40L-M90 (ATCC HB-12055) and M91 secreted by hybridoma hCD40L-M91 (ATCC HB-12056). M90 and M91 inhibit CD40-L binding to CD40 and the ability of triméric CD40-L and anti-Immunoglobulin M to induce
                                                                                                                                                                                                         TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
                                                                                                  099
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                                                                                                                                                                                                                                                                                                                                                                                                                                    antibody;
                                                           GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
                                                                                                               ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC
              GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGGTGAAGCCCAGCATTGGATCT
ACGITAAAAGAGAAGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGG
                                                                                                                                                                                                                                                                                                                                                                                                                                      monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                   CD40-L; CD40 receptor ligand; anti-human CD40-L monoclonal binding inhibitor; trimeric CD40-L; anti-Immunoglobulin M; peripheral blood B cell; proliferation inhibitor; ss:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Anti-human CD40-Ligand monoclonal antibodies
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                                                                                                                                                                                                                                                                                                                                         AAZ27524 standard; cDNA; 783 BP.
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91US-0805723.
92US-0969703.
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                                                                                                                                                                                                                                                                                                                                                                                                                   Mouse CD40-L coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94US-0249189
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23-OCT-1992;
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DB 20; Length 783;

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Score

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Query Match

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         Gaps
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transmembrane; region; intracellular; soluble; activity; B cell;
proliferation; induction; antibody; secretion; IgE; agonist;
                                                                                            TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA
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Similarity
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proliferation;
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This sequence encodes a murine CD40-L polypeptide which binds to CD40. CD40-L is a type II membrane polypeptide which has an extracellular region at its C-terminus, a transmembrane region and an intracellular region at its N-terminus. A soluble form of CD40-L lacks the transmembrane domain. CD40-L activity is mediated by binding with CD40 and induces B cell proliferation and induction of antibody secretion, including IgE. Membrane bound CD40-L acts as a CD40 agonist and soluble CD40-L acts as a CD40 antibody secretion assay to detect cells expressing CD40-L
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New cytokine CD40-L as CD40 agonist and antagonist - is used for treating allergies, lupus, rheumatoid arthritis, graft-versus-host disease and insulin-dependent diabetes mellitus
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Pred. No. 2.2e-200;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                          Armitage RJ,
                                                                                                                                                                                                                                                         25-OCT-1991;
05-DEC-1991;
                                               musculus
                                                                                                                                                                                                       23-OCT-1992;
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Gaps 9 9 240

300

480

180

us-08-982-272-5.rng

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/note= "signal peptide from murine surfactant protein D"
                                      ACTGAAGCAAGCCAAGTGGATCCACAGAGTTGGCTTCTCTCTTTTGGCTTACTCAAACTC 780
                                                                                                                                   Fusion protein; CD40 ligand; tumour necrosis factor; TNF; collectin; pulmonary surfactant protein D. SPD; immunocompetant cell; cell antigenicity; vaccine adjuvant; chimera; SS.
                          540
                                                                   Nucleotide sequence of surfactant protein D fused to CD40 ligand.
aagaaaggatattataccatgaaaagcaacttggtaatgcttgaaaatgggaaacagctg
                   acggttaaaagagaaggactctattatgtctacactcaagtcaccttctgctctaategg
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                                                                                                                                                                                                           ВР
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P-PSDB; AAB84421.
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Producing tumor necrosis factor superfamily proteins as multimeric ligands fused onto collectin molecules e.g. pulmonary surfactant protein D, useful as vaccine adjuvants against infectious agents and

Claim 1; Page 65-68; 74pp; English.

tumours

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The present sequence encodes a fusion protein comprising the mature murine surfactant protein D (including hub region, collagemous portion, and neck) fused to the murine CD40 ligand extracellular region (including stalk). The specification describes a method for constructing stable constructing stable constructing stable constructing stable constructing fusion proteins of the difficult to express tumor necrosis construction proteins of the invention of the difficult protein packbones such as collectin molecules e.g. pulmonary surfactant protein D (SPD). The fusion proteins of the invention are useful for stimulating immune response in proteins of the invention are useful for stimulating immune response in useful for increasing antigensity of cells such as tumor cells or human useful for simulating stimulating immune response in immunodeficiency virus (HIV) positive cells. They are also useful as a immunodeficiency virus (HIV) positive cells. They are also useful as a immunodeficiency virus in the soluble fusion protein makes them cells in the large size of the soluble fusion protein makes them cells is such as vaccine adjuvant or tumor immunotherapy agent, injected locally used as a vaccine adjuvant or tumor immunotherapy agent, injected locally conteins present new possibilities for the expression of highly active, multimeric, soluble TNFSF members. CD40L was a powerful stimulant for macrophages and dendritic cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 645.2; DB 22; Length 1477;
Pred. No. 1e-169;
0; Mismatches 3; Indels 0;
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Local Similarity 99.5%;
hes 647; Conservative (
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Thu May

18-MAR-1996

AAT05762;

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RESULT AAT0576

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neoplasia;
361 ATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human, alteration; immunoreactivity; human cell;
accessory molecule ligand; AML; gene therapy; treatment;
autoimmune disorder; rheumatoid arthritis; vaccine; chime
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Exemplary nucleotide sequence of a chimeric accessory
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Chimeric - Mus sp.
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tga 795
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09-DEC-1996;
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                                                                                                                                                    High density membrane-bound CD40 ligand; B-lymphocyte; B-Cell; differentiation; proliferation; baculovirus; Spodoptera frugiperda; Sf9; insect cell culture; tumour necrosis factor receptor; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 642.2; DB 10;
. No. 5.6e-169;
.-has 88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             High density membrane bound CD40 ligand - for proliferation of B cells in vitro or in vivo,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
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88.8%;
                              AAT05762 standard; DNA; 818
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                                                                                                                         Murine CD40 ligand DNA
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differentiated cells
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P-PSDB; AAR85485.
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Best Local Similarity
Matches 695; Conserv
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28-APR-1994;

Castle BE,

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28-APR-1995;

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where nucelotides encoding human domain III have been replaced with nucleotide encoding human domain III have been replaced with nucleotide encoding mouse domain III. The sequence is used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells. Vectors containing the AML genes can be used in gene therapy for treating neoplasia or autoimmune disorders such as rheumatoid arthritis. They can also be used for vaccination to produce immunity against a virus cell,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     009
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            altering immunoreactivity of cells, particularly for treatment of neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
                                                                                                                                                                                                                                                                                                               ;
                                                                                                                                                                                              accination to produce immunity against a virus cell, protein, fungus or neoplasia.
                                                                                                                                                                                                                                                                                Length 783;
                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                         Sequence 783 BP; 252 A; 167 C; 166 G; 198 T; 0 other;
                                                                                                                                                                                                                                                                                                            90;
                                                                                                                                                                                                                                                                                   Score 639; DB 19;
Pred. No. 4.3e-168;
0; Mismatches 90;
                                              Disclosure; Page 116; 167pp; English
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Local Similarity 88.5%;
Les 693; Conservative
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This sequence encodes the mouse trimeric CD40 receptor ligand (CD40-L). The invention relates to anti-human CD40-L monoclonal antibodies M90 secreted by hybridoma hCD40L-M90 (ATCC HB-12055) and M91 secreted by hybridoma hCD40L-M91 (ATCC HB-12056). M90 and M91 inhibit CD40-L binding to CD40 and the ability of trimeric CD40-L and anti-Immunoglobulin M to induce proliferation of peripheral blood B cells.
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antibody;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 878;
                                                                                                                                                                                                                         CD40-L; CD40 receptor ligand; anti-human CD40-L monoclonal binding inhibitor; trimeric CD40-L; anti-Immunoglobulin M; peripheral blood B cell; proliferation inhibitor; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 878 BP; 277 A; 178 C; 203 G; 220 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 631.4; DB 20
pred. No. 5.9e-166;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Anti-human CD40-Ligand monoclonal antibodies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Armitage RJ;
                                                                                                                                                                                                         Mouse trimeric CD40-L coding sequence
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                                                                                                                                 ВР
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99.8%;
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91US-0805723.
92US-0969703.
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                                                                                                                                                                                    (first entry)
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P-PSDB; AAX39941.
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23-OCT-1992;
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                                                                                                                                                                                                                                                                                                                                                           24-MAY-1994;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CD40 ligand; alteration; immunoreactivity; human cell; accessory molecule ligand; AML; gene therapy; treatment; neoplasia; autoimmune disorder; rheumatoid arthritis; vaccine; chimera; ss.
                                                                                                                                           ttggtaatgcttgaaaatgggaaacagctgacggttaaaaagaaggactctattatgtc
                                                                                                                                                                                               gaaatgcaaagaggtgatgaggatcctcaaattgcagcacacgttgtaagcgaagccaac
                                                  TACACTCAAGTCACCTTCTGCTCTAATCGGGAGCCTTCGAGTCAACGCCCATTCATCGTC
                                                                                                                                                                                                                                                     GGCCTCTGGCTGAAGCCCAGCATTGGATCTGAGAGAATCTTACTCAAGGCGGCAAATACC
                                                                                                                                                                                                                                                                                                                          CACAGTTCCTCCCAGCTTTGCGAGCAGCAGTCTTTGTTCACTTGGGCGGAGTGTTTGAATTA
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                                AGTAATGCAGCATCCGTTCTACAGTGGGCCAAGAAAGGATATTATACCATGAAAAGCAAC
                                                                                                      TTGGTAATGCTTGAAAATGGGAAACAGCTGACGGTTAAAAGAGAAGGACTCTATTATGTC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Exemplary CD40 ligand gene used in the course of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Vectors containing accessory molecule lial altering immunoreactivity of cells, part neoplasia or autoimmune disorders, e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GGCTTCTCATCTTTGGCTTACTCAAACTCTGA
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Chimeric - Homo sapiens
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cytoplasmic domain (Domain I). The sequence is used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells. Vectors containing the AML genes can be used in gene therapy for treating neoplasia or autoimmune disorders such as rheumatoid arthritis. They can also be used for vaccination to produce immunity against a virus cell, bacteria,
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human CD40 ligand
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                                                                                                                                                                                                                                                                                   Length 786;
                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                        G; 201 T; 0 other;
                                                                                                                                                                                                                                                                                Score 592.4; DB 19;
Pred. No. 4:2e-155;
); Mismatches 111; 1
is operatively linked to the
                                                                                                                                                                                                                                                                                                                              0; Mismatches
                                                                                                                                                                                                                     Sequence 786 BP; 250 A; 168 C; 167
                                                                                                                                                                                neoplasia.
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85.5%;
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CAGTCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAAC 717
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                                                                                                             CAAATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGG
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ottttgctgtgtatcttcatagaaggttggacaagatagaagatgtgaagatcttcat
                                        GAAGATTTTGTATTCATAAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CD40 ligand; alteration; immunoreactivity; human cell; accessory molecule ligand; AML; gene therapy; treatment; autoimmune disorder; rheumatoid arthritis; vaccine; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CD40 ligand gene used in the course of the invention
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09-DEC-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CD40-L was ligated to a synthetic oligonucleotide sequence coding for a leader peptide, a 33 amino acid leucine zipper sequence (AAR53968) and the Flag (RTW) linker sequence. Cells expressing the fusion construct are grown to accumulate oligomeric, soluble CD40-L
                                                                                                                                                                                                                                                                                                                   /product human_CD40-L
/product= "nucleotides 148-783 code for the
/note= extracellular region (amino acids 50-261)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    in the supernatant, The leucine zipper sequence spontaneously trimerises in solution and fusion proteins comprising the sequence fused to a heterologous mammallan protein also form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 582.8; DB 15; Length 786;
Pred. No. 2e-152;
); Mismatches 117; Indels 3;
                                                                                                                                                                                Leucine zipper; trimerisation; trimeric CD40-L; fusion protein; hetero-oligomer; homo-oligomer; type II transmembrane protein; soluble CD40-L; tumour necrosis factor family; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Prepn. of soluble oligomeric mammalian proteins - using host cells to express a fusion protein comprising a leucine zipper domain and a heterologous mammalian protein
                                                                                                                                                          Human CD40-L type II transmembrane protein coding sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 786 BP; 250 A; 168 C; 168 G; 200 T; 0 other;
                                                                          AAQ63959 standard; cDNA to mRNA; 786 BP
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93US-0107353.
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Best Local Similarity 84.7
Matches 666; Conservative
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P-PSDB; AAR53969.
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13-AUG-1993;
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Cantwell M, Kipps TJ, Sharma S;

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Wetcors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatment of neoplasia or autoimmune disorders, e.g. rheumatoid arthritis

XX

Vectors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatment of neoplasia or autoimmune disorders, e.g. rheumatoid arthritis

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CC

Used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of human cells

CC used to exemplify the cells so that the AML is expressed on the surface of the cells. Vectors containing the AML is expressed on the surface of the cells. Vectors containing the AML genes can be used in gene cof therapy for treating neoplasia or autoimmune disorders such as rheumatoid arthritis. They can also be used for vaccination to produce immunity a virus cell, bacteria, protein, fungus or neoplasia.

XX

Sequence 786 BP; 250 A; 168 C; 168 G; 200 T; 0 other;
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240
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                                                                                              ATGAAAATTTTTATGTATTTACTTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
                                                                                                         CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180
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                                                             Gaps
                                                ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
                                                                                                                                                      TCTGAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGGCTTGCGAGCAG
                                                                                                                                                                                          GAAGATTTTGTATTCATAAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
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                          3;
  Length 786;
                           Indels
  DB 19;
Score 582.8; DB 19;
Pred. No. 2e-152;
0; Mismatches 117;
Query Match
Best Local Similarity 84.7%;
Matches 666; Conservative
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completed: May 30, 2002, 03:03:23 ne: 11252 sec

Search com

GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd.

OM nucleic - nucleic search, using sw model

May 30, 2002, 05:22:53 ; Search time 8356.06 Seconds (without alignments) 2027.286 Million cell updates/sec Run on:

US-08-982-272-5 783 Title: Perfect score: Sequence:

1 ATGATCGAAACATACAACCA.....TTGGCTTACTCAAACTCTGA 783

Scoring table:

IDENTITY_NUC Gapox 1.0

43959072 Total number of hits satisfying chosen parameters:

21979536 seqs, 10817449327 residues

Searched:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

'pna/US098A_COMB.seq 'pna/US098B_COMB.seq /pna/US099A_COMB.seq /pna/US099B_COMB.seq pna/US095B_COMB.sec 'pna/US098C_COMB.seq 2/pna/US085_COMB.seq: ptodata/2/pna/US089_COMB.seq ptodata/2/pna/US090_COMB.seq ptodata/2/pna/US091_COMB.seq ptodata/2/pna/US092_COMB.seq ptodata/2/pna/US088_COMB.sec ptodata/2/pna/US082_COMB.seq ptodata/2/pna/US083_COMB.seq ptodata/2/pna/US084_COMB.seq /2/pna/US087_COMB.sec /pna/US096A_COMB. 'pna/US096B_COMB. pna/US097B_COMB. COMB. Pending_Patents_NA_Main:*

n2_6/ptodata/2/pna/US6005_COMB.seq:* n2_6/ptodata/2/pna/US6006_COMB.seq:*	2_6/ptodata/2/pna/US6007_	2_6/ptodata/2/pna/US6008_	2_6/ptodata/2/pna/US6009_	2_6/ptodata/2/pna/US6010_	2_6/ptodata/2/pna/US6011_	2_6/ptodata/2/pna/US6012_	2_6/ptodata/2/pna/US6013_	2_6/ptodata/2/pna/US6014_	2_6/ptodata/2/pna/US6015_	2_6/ptodata/2/pna/US6016_	2_6/ptodata/2/pna/US6017_C	2_6/ptodata/2/pna/US6018_	2_6/ptodata/2/pna/US6019_C	2_6/ptodata/2/pna/US6020_C	2_6/ptodata/2/pna/US6021_C	2_6/ptodata/2/pna/US6022_C	2_6/ptodata/2/pna/US6023_C	2_6/ptodata/2/pna/US6024_C	2_6/ptodata/2/pna/US6025_C	n2_6/ptodata/2/pna/US6026_	n2_6/ptodata/2/pna/US6027_	n2_6/ptodata/2/pna/US6028_C	n2_6/ptodata/2/pna/US6029_C	2_6/ptodata/2/pna/US6030_C	2_6/ptodata/2/pna/US	2_6/ptodata/2/pna/US6032_	n2_6/ptodata/2/pna/US6033_	n2_6/ptodata/2/pna/US6034_COMB.se	n2_6/ptodata/2/pna/US6035_COMB.s	C	
20	'N	: /cgn2_	N.	: /cgn2_	: /cgn2_			2	: /cgn2	: /cgn2	: /cgn2	: /cgn2	: /cgn2	: /cdu	: /cgn2,	: /cgn2,	: /cdu	: /cgn2,	: /cdu	: /cdn	: /cgn2	: /cdu	: /cdu	: /cdu	: /cdu	: /cdu	: /cdu	: /cdu	gn	C	
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result	ar con	Query	Query Match Length	a	£	Description
:	31000		Total cu	3 :		
1	783	100.0	783	13	US-08-982-272-5	Sequence 5, Appli
ŝ	773.4	98.8	783	13	US-08-982-272-3	Sequence 3, Appli
٣	9.094	97.1	783	٣	US-07-969-703B-1	1, 7
4	9.09/	97.1	783	13	US-08-982-272-2	e 2,
S	9.09/	97.1	818	9	US-08-234-580-1	1, 7
9	759	96.9	783	ო	US-07-969-703A-1	Sequence 1, Appli
7	759	6.96	783	ស	US-08-107-353-5	Sequence 5, Appli
œ	759	6.96	783	8	-477-733	Sequence 1, Appli
σ	759	96.9	783	8	US-08-484-624-1	1,
10	759	6.96	783	11	US-08-770-981-1	e 1,
11.	759	6.96	783	14	US-09-088-913-1	'n
12	759	6.96	783	17	US-09-322-021-1	٦,
13	759	6.96	783	17	US-09-322-021A-1	1,
14	759	96.9	783	17	US-09-365-940-1	ı,
15	759	96.9	783	17	-365-940	1,
16	759	96.9	783	17	US-09-392-618-1	Sequence 1, Appli
17	759	6.96	783	17		ı,
18	759	96.9	1209	75	US-60-360-207-	Sequence 9611, Ap
19	759	96.9	1250	37		Sequence 3, Appli
20	757.6	96.8	780	m	US-07-783-707A-1	'n,
21	757.6	96.8	780	m	US-07-805-723-1	Sequence 1, Appli
22	645.2	82.4	1477	18	-223-	
23	639	81.6	783	13	US-08-982-272-20	•
24	631.4	80.6	878	œ	-477	~
25	631.4	90.6	878	ထ	US-08-484-624-22	Seguence 22, Appl
26	631.4	90.6	878	11	US-08-770-981-22	e 22,
27	631.4	80.6	878	14	US-09-088-913-22.	22,
28	631.4	80.6	878	17	US-09-322-021-22	22,
59	631.4	90.6	878	17	US-09-322-021A-22	
30	631.4	90.6	878	17	US-09-365-940-22	22,
31	631.4	90.6	878	17	US-09-365-940A-22	22,

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721 ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC 780
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                                                                                         1 ATGAICGAAACAIACAAACTICICCCCGAICIGCGGCCACIGGACIGCCCAICAGC 60
                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG
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                                                                                                                                                                                                                                                                                                                                                                                                                                               ACGGTTAAAAGGAGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGG
                                                                                                                                                                                 GAAGAITITGTATTCATAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
                                                                                                                                                                                                                                                241 TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA
                                                                                                                                                                                                                                                                                      AACAAAGAAGAAGAAAAAGAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAA
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                                                                                                                                                                                                                                                                                                                                                       421 AAGAÀAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG
                                                                                                                                                                                             241 TTGCTGAACTGTGAGGGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA
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CONTAINING ACCESSORY
MOLECULE LIGAND GENES AND
THEIR USE FOR IMMUNO-
THEIR USE FOR IMMUNO-
MODULATION AND TREATMENT OF
MALIGNANCIES
  ö
    Mismatches
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Sharma, Sanjai
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        Conservative
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US-08-982-272-3
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Sequence 2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 783; DB 13;
Pred. No. 2.8e-185;
                                                                                                                                                                                                                                                                                                                                                                                                  THEIR USE FOR IMMUNO-
MODULATION AND TREATMENT OF
                                                                                                                                                                                                                                                                                                                                                                         CONTAINING ACCESSORY MOLECULE LIGAND GENES AND
              US-09-392-618-22
US-09-399-106-22
US-08-982-272-4
                                                    US-08-107-353-3
US-08-982-272-1
PCT-US94-00786-7
PCT-US94-14767-1
PCT-US97-11956-1
                                                                                                                                                                                                                                                                                                                                                           NOVEL EXPRESSION VECTORS
                                                                                                                                     US-07-969-703A-11
US-07-969-703B-11
                                                                                                                        US-07-805-723-10
                                                                                                                                                              -08-009-258-7
                                                                                                                                                                          US-08-172-664-1
US-08-215-862-1
                                                                                                                                                                                                                              ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SYSTEM: IBM P.C. DOS 5.0
FastSeq Version 2.0
ICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US/08/982, 272
                                                                                                                                                                                                                                                                                                                                                                                                                             MALIGNANCIES
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                                                                                                                                                   US-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              633 West Fifth Street
                                                                                                                                                                                                                                                                                           Sequence 5, Application US/08982272 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: 3.5" Diskette, MEDIUM TYPE: 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Guise, Jeffrey W. REGISTRATION NUMBER: 34,613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 23
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 12/9/96 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (213) 955-0440
                                                                                                                                                                                                                                                                                                                                    Sharma, Sanjai
Cantwell, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 783 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                      Kipps, Thomas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Suite 4700
                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Los Angeles
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION
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cal Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
US-08-982-272-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE
                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
                                                                                                                                                                                                                                                                                   JS-08-982-272-5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE:
                                                                                                                                                                                                                                                                                                                                        APPLICANT:
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099 099 720

540

420

300

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REGISTRATION NUMBER: 34,693
REPERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 783 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 2065870606
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDALL
STREET: 51
CITY: SEATTLE
STATE: WASHINGTON
STATE: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: CDNA HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: MOUSE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IMMEDIATE SOURCE
CLONE: CD40-L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
NAME/KEY: CDS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           241 TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
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Pred. No. 7.2e-183;
0; Mismatches 6;
                                                                                                                                                                               OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: FastSeq Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/982,272
                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 231/003
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 3:
ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700 CITY: Los Angeles STATE: California
                                                                                                                                                                                                                                                                 CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/132145
                                                                                                                                  MEDIUM TYPE: 3.5" Diskette, MEDIUM TYPE: 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                               NAME: Guise, Jeffrey W. REGISTRATION NUMBER: 34,613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98.8%;
                                                                                                                                                                                                                                                                                                              FILING DATE: 12/9/96
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                   IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 98.8
Best Local Similarity 99.2
Matches 777; Conservative
                                                                                                                    COMPUTER READABLE FORM
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                                                                                 COUNTRY: U.S.A. ZIP: 90071-2066
                                                                                                                                                                                                                                                 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY:
US-08-982-272-3
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721 ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC
                                                              GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCCAGCATTGGATCT
                                                                                   GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGGCTTTGCGAGCAGCAG
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CURRENT APPLICATION DATA:
DELICATION NUMBER: US/07/969,703B
FILING DATE: 19921023
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
TITLE OF INVENTION: NOVEL CYTOKINE
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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APPLICANT: ARMITAGE, RICHARD
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CONTAINING ACCESSORY

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                                                                             1 ATGATCGAAACATACAACCAAACTICTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC:60
                                                                                          ACGGTTAAAAGGGAACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGG
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                                     Length 783;
                                                           Indels
                                       , DB 3;
.1e-179;
                                                ed. No. 1.1e
Mismatches
                                         Score 760.6;
                                                    Pred.
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GENERAL INFORMATION:
                                         97.1%;
                                                             Conservative
                                                      Similarity
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TGA 783
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                                         Query Match
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Matches 769
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g. ð Cantwell, Mark VENTION: NOVEL EXPRESSION VECTORS

TITLE OF INVENTION:

Sharma, Sanjai

APPLICANT: APPLICANT: APPLICANT:

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AACAAAGAAGAAAAAAAAAAAAAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAA 360
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; Pred. No. 1.1e-179;
0; Mismatches 14;
THEIR USE FOR IMMUNO-
MODULATION AND TREATMENT
                                                                                                                                                                                                                                                                                                                                  US/08/982,272
                                  MALIGNANCIES
                                                                                                                                                                                                                                                                               OPERATING SYSTEM: IBM P.C. DOS SOFTWARE: FASTSGQ VESTION 2.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/982,
                                                                                    ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                             3.5" Diskette,
1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Guise, Jeffrey W. REGISTRATION NUMBER: 34,613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 23
FELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/132
FILING DATE: 12/9/96
ATTORNEY/AGENT INFORMATION:
NAME: Guise Infferent
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98.2%;
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
TYPE: nucleic acid
cTRENDEDNESS: single
                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 98.2
Matches 769; Conservative
                                                                                                                                                                            COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
                                                                       CORRESPONDENCE ADDRESS
   TITLE OF INVENTION: TITLE OF INVENTION: MITTLE OF INVENTION: MUMBER OF SEQUENCES:
                                                                                                                        Suite 4700
                                                                                                                                           Los Angeles
California
                                                                                                                                                                                                                               MEDIUM TYPE:
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                                             GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
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                                                                              ACGCTTAAAAGAGAACGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGG
                                                                                                                           GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGATCT
                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: KEHRY, MERILYN R
APPLICANT: CASTLE, BRIAN E
TITLE OF INVENTION: METHODS FOR PROLIFERATING AND
TITLE OF INVENTION: DIFFERENTIATING B CELLS, AND USES THEREOF
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPENATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FLING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: STERN, KESSLER, GOLDSTEIN & FOX
STREET: 100 NEW YORK AVE. N.W. SUITE 600
CITY: WASHINGTON
STATE: D.C.
ZIP: 20005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1011.1030000/RAM
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28-APR-1994
                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/08234580 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MILLMAN, ROBERT A
REGISTRATION NUMBER: 36,217
REFERENCE/DOCKET NUMBER: 101.
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (202)371-2600
TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 818 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13..795
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TGA 783
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LOCATION:
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                                        Gaps
                                                                                                             72
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                                                                                              13 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC
                                                                                                                                                                     1. ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
                                                                                                                                                  61 ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG
                                                                                                                                                                                                                             CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGAGAAGTAAACCTTCAT
                                                                                                                                                                                                                                                                                                    GAAGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
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                                      Indels
 Score 760.6; DB 6;
Pred. No. 1.2e-179;
                                      Mismatches
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97.1%;
98.2%;
                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: ARMITAGE, APPLICANT: FANSLOW, W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION:
NUMBER OF SEQUENCES:
Query Match
Best Local Similarity
Matches 769; Conserv
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TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
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                                                                             541 GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGGTGAAGCCCAGCATTGGATCT
                                                                                                                                      601 GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCCAGCTTTGCGAGCAGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/107,353
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ADDRESSEE: Immunex Corporation
STREET: 51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 5, Application US/08107353 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
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CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
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STRANDEDNESS: single
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241 TIGCIGAACTGIGAGGAGAIGAGAAGGCAATITGAAGACCTIGICAAGAIATAACGITA 300
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Pred. No. 2.9e-179;
0; Mismatches 15; Indels
                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/969,703A
FILING DATE: 19921023
                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS.
           CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 28
                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION: 1ELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96.9%;
98.1%;
                                                                                                                                                                                                                        FILING DATE: 19921023
CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: PERKINS, PALTICIA A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 96.9
Best Local Similarity 98.1
Matches 768; Conservative
                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 783 base pairs TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                 TELEPHONE: Zuece
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                                                                        WASHINGTON
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                                                                        STATE: WP
COUNTRY:
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                                                                                                                                                                                                                                                        Length 783;
                                                                                                                                                                                                                                                                                               2.96-179;
                                                                                                                                                                                                                                                        DB, 5;
                                                                                                                                                                                                                                                                                                                                  0; Mismatches
                                                                                                                                                                                                                                            Ouery Match 96.9%; Score 759; Best Local Similarity 98.1%; Pred. No. 2 Matches 768; Conservative 0; Mismatch
CD40-L
                                                                  ; NAME/KEY:
; LOCATION:
US-08-107-353-5
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781 TGA 783
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61 ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
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APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 759; DB 8; 1
Pred. No. 2.9e-179;
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Microsoft Word for Apple,
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                                                                                        S: IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: December 5, 1991
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: OCTOBER 25, 1991
CLASSIFICATION: 435
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APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Perkins, Patricia A REGISTRATION NUMBER: 34,69
                                                                                                                                                                                                                                                       Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: May 24, 1994
CLASSIFICATION: 435
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Best Local Similarity 98.1%;
Matches 768; Conservative
                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 2065870606
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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nucleic acid
EDNESS: single
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                                                         NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS:
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                                                                                                                                   SEATTLE
WASHINGTON
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CLASSIFICATION:
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; LOCATION:
US-08-477-733A-1
                                                                                              ADDRESSEE:
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Sequence 1, Application US/08477733A GENERAL INFORMATION:

US-08-477-733A-1

APPLICANT: ARMITAGE, RICHARD

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98.1%; Pred. No. 2.9e-179;
iive 0; Mismatches 15;
                                                                                                                   NUMBER: 07/969,703
October 23, 1992
                                                                                                                                                                                                                                                           NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 28
                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION
                                               CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/249,
FILING DATE: May 24, 1994
CLASSIFICATION: 514
PRIOR APPLICATION: 514
PRIOR APPLICATION NUMBER: 07/969,
                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION
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nucleic acid
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2065870606
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                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE:
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                                                                                                                                                     PRIOR APPLICATION DATA: APPLICATION NUMBER:
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Best Local Similarity
Matches 768; Conserv
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                                                                                                                                           CLASSIFICATION:
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                                                                                                                                  FILING DATE:
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LOCATION:
US-08-484-624-1
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                                181 GAAGATTTTGTAITCATAAAAAGCTAAAGAGATGCAACAAAAGGAGAAGGATCTTTATCC 240
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APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GTBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
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COMPUTER: Apple Macintosh
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US-08-484-624-1
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Microsoft Word for Apple, version 5.1a
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FILING DATE: December 5, 1991
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CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
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PRIOR APPLICATION DATA
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                                                                                          AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG 480
                                                                                                                                                                                                              541 GAGCCTTCGAGTCAACGCCCATTCATCGTCGCCTCTGGCTGAAGCCCCAGCATTGGATCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: ARNITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 08/477,733
FILING DATE: 07-JUN-1995
FILING DATE: 08/249,189
FILING DATE: MAY 24, 1994
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/08770981 GENERAL INFORMATION:
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2IP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
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US-08-770-981-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 759; DB 11;
Pred. No. 2.9e-179;
0; Mismatches 15;
07/805,723
                                                                                       07/783,707
                                       CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,70
FILING DATE: October 25, 1991
                       December 5, 1991
                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: PERKINS, PALICIA A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 280
TELECHONE: 2065870430
TELEPHONE: 2065870606
                                                                                                                                                                                                   34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  96.9%;
98.1%;
                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID:NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ouery Match
Best Local Similarity 98.1
Matches 768; Conservative
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: CDNA HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: MOUSE
                                                                                                                                   CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ANTI-SENSE: NO
ORIGINAL SOURCE
                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; LOCATION:
US-08-770-981-1
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541 GAGCCTTCGAGTCAACGCCCATTCATCATCGCCCCTGGCTGAAGCCCAGCATGGATCT 600 541 GAGCCTTCGAGTCAACGCCCATTCATCATCGTCGGCCTCTGGCTGAAGCCCAGCGTGGATT 600
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                                                                                  TCTGTTCACTTGGGCCGAGGTTTGAAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
                                                                                                                                                    GAGAGAATCTTACTCAAGGCGGCAAATACCCCAGGTTCCTCCCAGCTTTGCGAGCAGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIA
APPLICANT: STRIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
CORRESPONDENCE: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    version 5.1a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: Apple Operating System 7.1 SOFTWARE: Microsoft Word for Apple, version CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA: 07/782
FILING DATE: 07/782
ATTORNO
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: October 23, 1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,725
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: October 25, 1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/09088913 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 2065870606
INFORMATION FOR SEQ ID NO:
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STREET: ST.
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721 ACTGAAGCAAGCGAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC 780
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                                                                                                                                             Length 783;
                                                                                                                                                                 Indels .
                                                                                                                                               96.9%; Score 759; DB 14;
98.1%; Pred. No. 2.9e-179;
iive 0; Mismatches 15;
                                                                                                                                                                     Conservative
      STRANDEDNESS: single TOPOLOGY: linear MOLECULE TYPE: CDNA HYPOTHETICAL: NO ANTI-SENSE: NO ORGANISM: SOURCE: IMMEDIATE SOURCE:
nucleic acid
                                                                                                                                                   Query Match
Best Local Similarity
Matches. 768; Conserv
                                                                                                     NAME/KEY:
                                                                                                              ;
US-09-088-913-1.
                                                                                    CLONE:
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241 TTGCTGAACTGTGAGGAGTGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         661 TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 721 ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC 780
                                                                                   61 ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
                                                                                                             61 ATGAAGATTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
                                                                                                                                                                         121 CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180
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1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
                                       1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                 AACAAAGAAGAGAAAAAAGAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: STNIVASAN, SUBHASHINI
APPLICANT: GIBSOM, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEE: IMMUNEX CORPORATION: 51 UNIVERSITY STREET
SEATTLE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Microsoft Word for Apple, version 5.1a CURENT APPLICATION DATA:
APPLICATION NUMBER: US/NO.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 759; DB 17;
Pred. No. 2.9e-179;
0; Mismatches 15;
                                                                                                                                                                  APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, WILLIAM
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, WARXLOU
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
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APPLICATION NUMBER: 07/783,70
FILING DATE: October 25, 1991
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: PERKINS, PARLIGIG A.
REGISTRATION NUMBER: 34,693
REFERRACE/DOCKET NUMBER: 2802
TELECOMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEPHONE: 2065870636
                                                                                                                           Sequence 1, Application US/09322021 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Op
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      96.98;
98.18;
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Best Local Similarity 98.1
Matches 768; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLEĆULE TYPE: CDNA
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: WASHINGTON
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IMMEDIATE SOURCE:
CLONE: CD40-L
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US-09-322-021-1
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WASHINGTON

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ADDRESSEE: STREET: CITY: S

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                                                                      aacaaagaagagaaaaaaagaaaacagctttgaaatgcaaagaggtgatgaggatcctcaa 360
                                                                                                                                                                                                                                                                                 601 GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
                                                                                                                                                                                                                                                                                              TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG
                                                                                                                                                                                                                                                                                                                                                    721 ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC
                                                                                                               AAGAAAGGATATTATACCATGAAAAGGAACTTGGTAATGCTTGAAAATGGGAAACACTG
                                                                                                                                                                                                                         541 GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGGAAGCCCAGCATTGGATCT
              361 ATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              System 7.5.5
Macintosh, version 6.0.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: MCGREW, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SYSTEM: Apple Operating MS Word for Apple Power
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APPLICATION NUMBER: US/09/365,940
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FANSLOW, WILLIAM
SPRIGGS, MELANIE
SRINIVASAN, SUBHASHINI
GIBSON, MARYLOU
MORRIS, ARVIA E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07/969,703
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          October 23, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1, Application US/09365940
GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple OF
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51 UNIVERSITY
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WASHINGTON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: SEATTLE STATE: WASHING
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         781 TGA 783
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US-09-365-940-1
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                                                                            SOFTWARE: Microsoft Word for Apple, version 5.1a SOFTWARE: Microsoft Word for Apple, version 5.1a CURRENT APPLICATION DATA: PAPLICATION NUMBER: US/09/322.071A FILTURE PRESENT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 759; DB 17;
Pred, No. 2.9e-179;
0; Mismatches 15;
                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
APPLICATION NUMBER: 07/805,723
CLASSIFICATION:
PRIOR APPLICATION DATA:
PAPPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/249,189
                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 34,693
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 280
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          96.9%;
nilarity 98.1%;
Conservative 0
                                                             E: Floppy disk
Apple Macintosh
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nucleic acid
                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy (
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Best Local Similarity
Matches 768; Conserv
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CLONE: CD40-L
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ORIGINAL SOURCE:
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                                   98101
                                                                                                                                                      FILING DATE
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LOCATION:
US-09-322-021A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pred. No.
                                                                                                                                       2802-D
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
CLASSIFICATION:
                                                           07/783,707
                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,70
FILING DATE: October 25, 1991
CLASSIFICATION:
                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFRENCE/CDCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                     96.9%;
98.1%;
                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                            LENGTH: 783 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                       STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                        2065870606
                                                                                                                                                                                                                                                                 MOLECULE TYPE: CDNA HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                  ORGANISM: MOUSE
IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                                        ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                            ANTI-SENSE: NO
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; LOCATION:
US-09-365-940-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                              768;
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Matches 76
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APPLICANT: ARMITAGE, Richard, J.
APPLICANT: FANSLOW, III, William, C.
APPLICANT: SPRIGGS, Melanide, K.
TITLE OF INVENTION: METHOD OF LEADING AN UNDESIRED IMMUNE RESPONSE BY BLOCKING TH
TITLE OF INVENTION: OF CD40 LIGAND TO CD40
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                                                                                                                        481 ACGGTTAAAAGAAGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGG
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                                541 GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCTCTGGCTGAAGCCCAGCATTGGATCT
                                                   GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
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Pred. No. 2.9e-179;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/365,940A CURRENT FILING DATE: 1999-08-02 PRIOR APPLICATION NUMBER: 08/349,189
PRIOR FILING DATE: 1994-05-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 07/969, 703
PRIOR FILING DATE: 1992-10-23
PRIOR PELLING DATE: 1991-10-23
PRIOR PELLING DATE: 1991-12-05
PRIOR APPLICATION NUMBER: 07/783,707
PRIOR APPLICATION NUMBER: 07/783,707
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 1, Application US/09365940A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98.18; Fig.
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Matches 768; Conservative
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NAME/KEY: CDS
LOCATION: (1)..(783
OTHER INFORMATION:
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781 TGA 783
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US-09-365-940A-1
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Search completed: May 30, 2002, 05:23:00 Job time: 18659 sec

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(without alignments)
1310.273 Million cell updates/sec
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                                                                                                                             May 30, 2002, 05:39:00; Search time 627.19 Seconds
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                                                 850503 seqs, 524770293 residues
                                                                                      OM nucleic - nucleic search, using sw model
                                                                                                                                                                                                                                                                                                                IDENTITY_NUC Gapop 10.0 , Gapext 1.0
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                                                                                                                                                                                                                        Title:
Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                 Searched:
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Pending_Patents_NA_New:*

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

/cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:* /cgn2_6/ptodata/1/pna/US06_NEW_COMB.seq: /cgn2_6/ptodata/1/pra/US07_NEW_COMB.seq. /cgn2_6/ptodata/1/pna/US08_NEW_COMB.seq. /cgn2_6/ptodata/1/pna/US09_NEW_COMB.seq. /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq. /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq. Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	Sequence 821, App		457,	Sequence 131, App	9, Ap	12	٠.,	П	П	Sequence 35082, A	_	Sequence 29, Appl	555,	9151,	Seguence 10646, A							Sequence 258659,	Sequence 239684,	Sequence 1404, Ap	Sequence 499, App	8655,
	QI	US-09-053-375B-821	US-09-053-375B-226	US-09-442-384B-457	US-09-911-904-131	US-09-875-453A-9	US-10-105-299-12515	US-09-830-706B-37	US-10-027-632-187249		US-10-027-632-35082	US-09-721-544-11180	US-10-071-192-29	US-09-789-189-555		US-10-105-299-10646	US-10-105-299-14274	US-10-105-694-1	US-10-105-695-1	US-10-106-014-1	US-10-027-632-199307	US-10-027-632-207524	US-10-027-632-258659	US-10-027-632-239684	US-10-106-698-1404	US-10-106-698-499	US-09-919-002-8655
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	Match Length DB	1250	1816	1816	508	2395	45698	99299	476	476	685	349	1758	341	57649	57649	57649	2430	2430	2430	1379	598	2580	626	712	1640	1665
% Query	Match	96.9	74.4	74.4	40.2	18:7	6.4	4 9	6.4	4.9	4.9	4.8	4.7	4.7	4.7	4.7.	4.7	4.6	4.6	4.6	4.6	4.6	4.6	4.5	4.5	4.5	45
	Score	759	582.8	582.8	315	146.2	38.4	38.4	38	38	. 38	37.6	37	36.6	36.6	36.6	36.6	36.2	36.2	36.2	36	35.8	35.8	35.6	35.6	35.6	35.6
Result	No.		7	m	4	'n	o o	c 7	œ	σ	c 10	11	12	c 13	14	15	16	17.	18	19	c 20	c 21	c 22	c 23	24	25	c 26

***************************************	Sequence 83187, A Sequence 46525, A Sequence 210428, Sequence 210394, Sequence 317028, Sequence 317029,	Sequence 228875, Sequence 233425,
US-10-027-632-219364 US-10-027-632-219365 US-10-027-632-219365 US-10-027-632-111808 US-10-027-632-111134 US-10-113-872-380 PCT-USO2-07826-362 US-10-097-340-362 US-10-097-632-85186	US-110-027-622-9518/ US-10-027-622-46525 US-10-027-632-210428 US-10-027-632-200394 US-10-027-632-317028 US-10-027-632-317029	US-10-027-632-228875 US-10-027-632-233425
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222 222 232 232 24 25 26	339 440 442 433	44 45
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ALIGNMENTS

RESULT 1 UG-09-053-375B-821 Sequence 821, Application US/09053375B SEQUENCE SEL, INFORMATION: APPLICANT: Chenchik, Alex APPLICANT: Bibliashvilli, Robert TILLE REFERENCE: CLON-006 CURRENT APPLICANTON: NUMBER: US/09/053,375B CURRENT FILING DATE: 1998-08-31 NUMBER OF SEQ ID NOS: 1543 SCOFTWARE: FastSEQ for Windows Version 4.0 SEQ ID NO 821 LENGTH: 1250 LENGTH: 1250 TYPE: DNA ORGANISM: MUS musculus US-09-053-375B-821	Ouery Match 96.9%; Score 759; DB 5; Length 1250; Best Local Similarity 98.1%; Pred. No. 2.8e-184; Matches 76%; Conservative 0; Mismatches 15; Indels 0; Gaps	1. ATGATCGAAACATACAACCATCCCCCATCTGCGGCCACTGGACTGCCCATCAGC 60	61 AIGAAAATTTTTAIGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGFG 120 	121 CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180 	181 GAAGATTTTGTATTCATAAAAAGCTAAAGACATGCAACAAGGAAGG	241 TTGCTGAACTGTGAGGAGATGAGAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
RESULT US-09-053 Sequence Sequence APPLIC TITLE FILE R CURREN CURREN CURREN CURREN SOFTWA SOFTWA SOFTWA TYPE: TYPE: TYPE: TYPE: TYPE: ORGAN	Query M Best Lo	Oy 1 Db 13	Qy 61 Db 73	Oy 121 Db 133	Oy 181 Db 193	Qy 241 Db · 253
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ATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC 420

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TYPE: DNA ORGANISM:
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                                                                                                                                                                   ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
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                                                                                                                   GAGCCTTCGAGTCAACGCCCATTCATCGTCGGCGTCTGGCTGAAGCCCAGCATTGGATCT 600 gagccttcgagtcaacgcccattcatcgtcggctctggctgaagcccagcagtggatct 612
                                                                                                                                                                                                               TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
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                                                                                                                                                                                                                            373 attgcagcacacgttgtaagcgaagccaacagtaatgcagcatccgttctacagtgggcc 432
                      480
                                 ACGITAAAAGAGAAGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAATCGG 540
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 582.8; DB 5;
Pred. No. 3.1e-139;
0; Mismatches 117;
                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Chenchik, Alex
APPLICANT: Bibilashvilli, Robert
TITLE OF INVENTION: Nucleic Acid Arrays
FILE REFERENCE: CLON-006
CURRENT APPLICATION NUMBER: US/09/053,375B
CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 1598
SEQ ID NO 226
SEQ ID NO 226
                                                                                                                                                                                                                                                                                                                                                                                                Sequence 226, Application US/09053375B
GENERAL INFORMATION:
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ilarity 84.7%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Homo sapiens
US-09-053-375B-226
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Matches 666;
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                                                                                                          CGGGAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGGCTGAAGCCCCAGCATTGGA 597
                                                                                                                                                                                                                                                                                                                                                                                          GTGACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAA
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340 aacaaagaggagacgaagaaagaaaacagctttgaaatgcaaaaaggtgatcagaatcct
                             CAAATTGCAGCACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCGGTTCTACAGTGG
                                               GCCAAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAAACAG
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Pred. No. 3.1e-139;
); Mismatches 117;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lukashev, Matvey
TITLE OF INVENTION: Hematology/Immunology Array
FILE REFERENCE: CLON-006CIPL5
FOREBREA PAPLICATION NUMBER: US/09/442,384B
CURRENT FILING DATE: 1999-11-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEO ID NOS: 830 SOFTWARE: FastSEO for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 457, Application US/09442384B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: 09/053,375 PRIOR FILING DATE: 1998-03-31
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84.78;
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US-09-442-384B-457
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Best Local Simi
Matches 666;
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TTACAAGCTGGTGCTTCTGTGTTTGTCAACGTGACCTGAAGCCAAGCCAAGTGATCCACAGA 747
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                                                                                                                          388 AACAGTAATGCAGCATCCGTTCTACAGTGGGCCAAGAAAGGATATTATACCATGAAAAGC
                                                                                                                                                                                                                                       GTCTACACTCAAGTCACCTTCTGCTCTAATCGGGAGCCTTCGAGTCAACGCCCATTCATC
                                                                                                                                                                                                                                                                                                                                                  628 ACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAGTCTGTTCACTTGGGCGGAGTGTTTGAA
              748 GTTGGCTTCTCATCTTTTGGCTTACTC
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Lim, Moon Young
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Tam, Albert W.
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Thomas, Rita L.
Kongpachith, Ana
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US-09-875-453A-9
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US-09-875-453A-9
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LENGTH: 2395
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APPLICANT:
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 ctttttgctgtgtatcttcatagaaggttggacaagatagaagatgaaaggaatcttcat
                                       CAAATTGCAGCACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGG
                                                                                                                                                                                                                                                   GCCAAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAG
                                                                                                                                                                                                                                                                CTGACGGTTAAAAGAGAAGGACTCTATTATGTCTACACTCAAGTCACCTTCTGCTCTAAT
                                                                                                                                                                                                                                                                                                                                                             CGGGAGCCTTCGAGTCAACGCCCATTCATCGTCGGCCTCTGGCTGAAGCCCAGCATTGGA
                                                                                                                                                                                                                                                                                                                                                                                                                  TCTGAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CAGTCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTTGTCAAC
                           GAAGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAAGGAGGAGGATCTTTATCC
                                                                                                                                       AACAAAGAAGA - - - GAAAAAAGAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 5e-71;
0; Mismatches 120;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Neft, Robin Elleen
APPLICANT: Neft, Robin Elleen
TITLE OF INVENTION: CANINE TOXICITY GENES
FILE REFERENCE: 400742000200
CURRENT APPLICATION NUMBER: US/09/911,904
CURRENT FILING DATE: 2002-04-09
PRIOR FILING DATE: 2000-07-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 315;
Pred. No. 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQ ID NOS: 386
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 131
LENGTH: 508
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 131, Application US/09911904
GENERAL INFORMATION:
APPLICANT: Farr, Spencer B.
APPLICANT: Pickett, Gavin G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     40.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TYPE: DNA
; ORGANISM: Canis familiaris
US-09-911-904-131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ouery Match 40.2
Best Local Similarity 76.3
Matches 387; Conservative
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Pred. No. 1e-27;
0; Mismatches 23;
                                                                                                                                                                                                                                                                                   APPLICANT: Bruice, Thomas W.
TITLE OF INVENTION: PROMOTERS FOR REGULATER
FILE REFERENCE: 54600-8135.USO0
CURRENT APPLICATION UNBER: US/09/875,453A
CURRENT FILING DATE: 2002-05-06
                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 246
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: US 60/209,549 PRIOR FILING DATE: 2000-06-06
Sequence 9, Application US/09875453A GENERAL INFORMATION:
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87.48;
                                                                                                                                   Michelotti, Emil F
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Best Local Similarity 87.4
Matches 160; Conservative
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                                                                                                                                                                                                                                                                                         248 ACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAG 307
                                                                                                                                                                                                                    Tength .66566;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 187249, Application US/10027632
GENERAL INFORMATION:
APPLICANT: Wang, David
TITLE OF INVENTION: Identification and Mapping of Sing-
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
                                                                                                                                                                                                                                                                                                                                                                       308 AAGAGAAAAAGAAAACAGCTTTGAAATGCAAAGAGGTGATGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches 100;
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Pred. No. 2.6;
                                                                                                                                                                                                                                     1 Similarity 60.6%, Pred. No. 9.4; 63; Conservative 0; Mismatches
                                                                                                                                                                                                                4.9%; Score 38.4;
60.6%; Pred. No. 9.4
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FastSEQ for Windows Version 4.0
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US 60/156,358
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PRIOR APPLICATION NUMBER: US 60/198,676
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LICATION NUMBER: US 60/167,363
                                                                                                                                      OTHER INFORMATION: n is unknown 09-830-706B-37
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                        ORGANISM: Homo sapiens
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Matches 98; Conserv
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PatentIn
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US-10-027-632-187249
                                                                                                              NAME/KEY: unsure
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Matches 63
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SOFTWARE:
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217 AACAAAGGAGAAGGATCTTTATCCTTGCTGAACTGTGAGGAGGATGAGAAGGCAATTTGAA 276
                                                                          ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
                                                                                                                                                      CITITICCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180
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TITLE OF INVENTION: Human Secreted Proteins
FILER REFERENCE: PS950
CURRENT APPLICATION NUMBER: US/10/105,299
CURRENT FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 15197
Prior Application removed - See File Wrapper or Palm
SOFWARE: Patentin Ver. 2.0
SEQ ID NO 12515
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APPLICANT: TAMAI, KATSUVUKI
TITLE OF INVENTION: THIOREDOXIN REDUCTASE II
FILE REFERENCE: 55865-71965
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PRIOR FILING DATE: 1999-10-28
PRÎOR APPLICATION NUMBER: JP 1998-310422
                                                                                                                                                                                                                                                                                                                                                               Sequence 12515, Application US/10105299 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Homo sapiens
US-10-105-299-12515
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Best Local Similarity
Matches 93; Conserv
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US-09-830-706B-37/c
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                                                                                                                                                                             Sequence 187250, Application US/10027632
GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide TITLE OF INVENTION: Polymorphisms in the Human Genome FILE REFERENCE: 108927.129

CURRENT APPLICATION NUMBER: US/10/027,632

CURRENT FILING DATE: 2002-04-30
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Pred. No. 2.6;
0; Mismatches 100; Indels
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PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/10/027,632 CURRENT FILING DATE: 2002-04-30
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PRIOR FILING DATE: 2000-07-12
PRIOR PELLING DATE: 2000-04-20
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
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PRIOR PAPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
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PRIOR APPLICATION NUMBER: US 60/185,218
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PRIOR APPLICATION NUMBER: US 60/167,363
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449 ACTIGGTAATGCTTGAAA 466
                                                     219 aatggagaataataaaaa 236
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Best Local Similarity 49.55
Matches 98; Conservative
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                                                                                                                                                       US-10-027-632-187250
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US-10-027-632-187250
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318 AAAATGAAAATCATGTTAAAAACTTGTGCAAAAGTGTTGTGATATTAAAATATAATACTAA 259
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Pred. No. 2.9;
0; Mismatches 100;
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR PILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1909-11-23
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR PILING DATE: 1999-11-23
PRIOR PILING DATE: 1999-10-28
PRIOR APPLICATION NUMBER: US 60/15,358
PRIOR PILING DATE: 1999-09-28
PRIOR PILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-09
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SOFTWARE: FastSEQ for Windows Version 4.0
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; OTHER INFORMATION: n = A,T,C or G
US-10-027-632-35082
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Palencia, Servando
Raisi, Fariba
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Best Local Similarity 49.5%;
Matches 98; Conservative
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Dickson, Mark
Drake, Jim
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Faulkner, Brandy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                García, Veronica
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US-09-721-544-11180
                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                    SEQ ID NO 35082
                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
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Benjamin

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LOCATION: 1136
SEQUENCE DESCRIPTION: SEQ ID NO: 29:
                                                          LENGIH: 1758 base pairs
TELEFAX: 303/863-0223
                                                                           TYPE: nucleic acid
STRANDEDNESS: single
               INFORMATION FOR SEQ ID NO: 29: SEQUENCE CHARACTERISTICS:
                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: CDNA
                                                                                                                                                                                  NAME/KEY:
                                                                                                                                                                                                       LOCATION:
                                                                                                                                                                                                                                                NAME/KEY:
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                                                                                                                                                                                                                               FEATURE
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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  316 AAAGAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAAATTGCAGCACACGTT 375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                    TITLE OF INVENTION: Novel Nucleic Acid Sequences Obtained IIILE OF INVENTION: From a CDNA Library of Fetal Liver-Spleen
                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 349;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                54; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NOVEL ECTOPARASITE SALIVA APPARATUS TO COLLECT SUCH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Weber, Eric R.
TITLE OF INVENTION: NOVEL ECTOPARASITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: SHERIDAN ROSS P.C.
STREET: 1560 BROADWAY, SUITE 1200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/071,192
FILING DATE: 07-Feb-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: 09/171,156
                                                                                                       CURRENT APPLICATION NUMBER: US/09/721,544 CURRENT FILING DATE: 2000-11-21
                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 24489
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: <Unknown>
                                                                                                                                                                       PRIOR APPLICATION NUMBER: 09/515,128
                                                                                                                                                                                             PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: 09/034,341
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-10-071-192-29
Sequence 29, Application US/10071192
GENERAL INFORMATION:
GENERAL SHOLICANT: Hunter, Shirley Wu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            JTER READABLE FORM:
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Best Local Similarity 56.5%;
Matches 70; Conservative
                                                                                                                                                                                                                                        PRIOR FILING DATE: 1998-02-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCES:
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                                                     Yim, Kenneth
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LENGTH: 349
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203 AGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCCTTGCTGAACTGTGAGGAGATGA 262
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                                                                                                                                                                                                       1103 AAGAAACTAAAGGAGCTCAACGATTGTTGAAGAWGGTTAACAAGATGATAACGAAAATGG 1162
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Length 1758;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Lelias, Jean-Michel TITLE OF INVENTION: Human Polynucleotides and Polypeptides TITLE OF ENERRICE: 25436/1720 CURRENT APPLICATION WHERE: US/09/789,189 CURRENT FILING DATE: 2001-02-20
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                                              1; Mismatches, 121;
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0; Mismatches
  Score 37;
Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 555, Application US/09789189
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      4.78;
                          Best Local Similarity 47.2
Matches 109; Conservative
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Matches 78; Conserv
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158 AGGTTGTGAAAAATAAAAGATGTAATG 132

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Search completed: May 30, 2002, 05:39:58 Job time: 16772 sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   161 AAGAGGAAGTAAACCTTCATGAAGATTTTGTATTCATAAAAAGCTAAAAGAGATGCAACA 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 221 AAGGAGAAGGATCTTTATCCTTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACC 280
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10646, Application US/10105299; GENERAL INFORMATION:
TITLE OF INVENTION: Human Secreted Proteins; FILE REFERENCE: PS950
CURRENT APPLICATION NUMBER: US/10/105,299; CURRENT APPLICATION NUMBER: US/10/105,299; NUMBER OF SEQ ID NOS: 15197; Prior Application removed - See File Wrapper or Palm; SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10646
LENGTH: 57649
                                                                             APPLICANT: Rosen, et. al
TITLE OF INVENTION: Human Secreted Proteins
FILE REFERRNCE: PS950
CURRENT APPLICATION UNMBER: US/10/105,299
CURRENT FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 15197
Prior Application removed - See File Wrapper or Palm
SEQ ID NO 9151
LENGTH: 57649
                                                                                                                                                                                                                                                                                                                                                                                                  Score 36.6; DB.6; I Pred. No. 26; 0; Mismatches 64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 36.6; DB 6;
Pred. No. 26;
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RESULT 14
US-10-105-299-9151
Sequence 9151, Application US/10105299
GENERAL INFORMATION
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Best Local Similarity 54.0%;
Matches 75; Conservative (
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Conservative
                                                                                                                                                                                                                                                                                              ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-105-299-9151
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; ORGANISM: Homo sapiens
US-10-105-299-10646
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Best Local Similarity
Matches 75; Conserva
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US-10-105-299-10646
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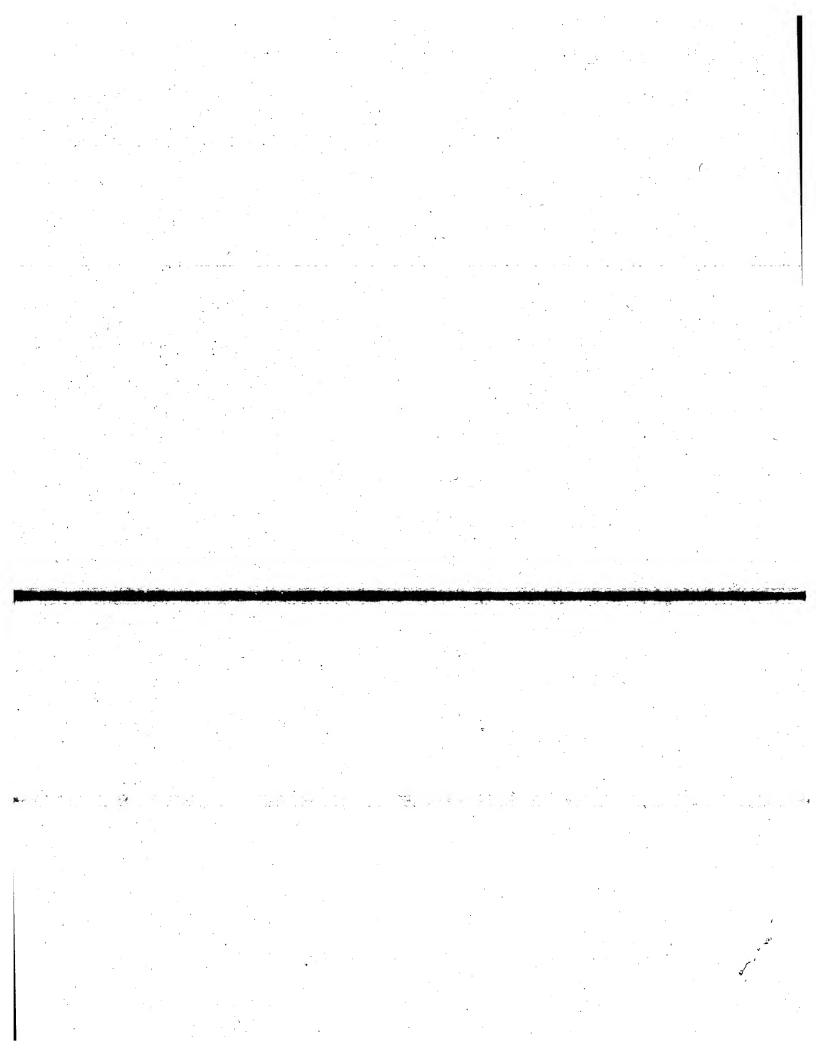
TTGTCAAGGATATAACGTT 299

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May 30, 2002, 02:51:32 ; Search time 139.75 Seconds (without alignments) 1376.251 Million cell updates/sec
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1 ATGATCGAAACATACAACCA......TTGGCTTACTCAAACTCTGA 783
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                         383533 seqs, 122816752 residues
                                                                                   OM nucleic - nucleic search, using sw model
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Perfect score:
Sequence:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

1: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/2/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/2/ina/6A_COMB.seq:*
4: /cgn2_6/ptodata/2/ina/6B_COMB.seq:*
5: /cgn2_6/ptodata/2/ina/PCTUS_COMB.seq:*
6: /cgn2_6/ptodata/2/ina/PcTUS_COMB.seq:*

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Issued_Patents_NA:

Database :

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SUMMARIES		-431-055-1	-858-197-1	-446-922-5	-249-189-1	-484-624A-1	-477-733B-1	-088-913A-1	US-08-769-819-1	US-08-770-974-1	PCT-US93-10034-5	US-08-249-189-22	A-2	US-08-477-733B-22	US-09-088-913A-22	-769-819-22	US-08-770-974-22	US-08-446-922-3	PCT-US93-10034-3	US-07-940-605A-1	US-08-184-422-7	US-08-360-923A-1	US-08-431-055-3	US-08-690-096-1	US-08-249-189-11	-484-624A-1	-477-733B-1	-08-763-995-1
SUS	ΞĎ	US-08-	ns-08	-80-Sn	ns-08	-80-Sn	ns-08	.60-SD	US-08	.80-SD	PCT-US	ns-08	ns-08	ns-08	0S-09	ns-08	ns-08	ns-08	PCT-US	US-07	ns-08	0S-08	ns-08	0S-08	ns-08	us-08	0S-08	0S-08
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ď	Query Match	97.1	97.1	6.96	6.96	6.96	6.96	6.96	6.96	6.96	6.96	90.6	90.6	90.6	90.6	9.08	90.6	74.4	74.4	74.4	74.4	74.4	74.4	74.4	74.4	74.4	74.4	74.4
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97.1%; Score 760.6; DB 1; Length 818; 98.2%; Pred. No. 1.6e-220;

Query Match Best Local Similarity

CDS 13..795

; NAME/KEY; ; LOCATION: US-08-431-055-1

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9-088-913A-11 5-589-71B-7 7-70-974-11 8-770-974-11 8-858-197-3 8-249-189-15 8-477-733B-15 9-769-819-15 8-770-974-15 8-770-974-15 8-770-974-15 8-770-974-15 8-770-974-15 8-770-974-15 8-770-974-15 8-484-624A-20 8-484-624A-20 8-484-624A-20 8-484-624A-20 8-488-9189-20 8-488-9189-20 8-488-9189-20 8-488-9189-20 8-488-9189-20 8-488-9189-20 8-488-9189-20 8-488-9189-20 8-488-9189-20 8-769-819-20	ALIGNMENTS	ROLIFERATIONG B CELLS STEIN & FOURTHE 600 SUITE 600 55 80
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80 80 80 80 80 80 80 80 80 80 80 80 80 8		REGULT 1 Sequence 1, Applicat Sequence 1, Applicat Sequence 1, Applicat Batent No. 5817516 GENERAL INFORMATION APPLICANT: CASTL TITLE OF INVENTIO TITLE OF INVENTIO TITLE OF INVENTIO TITLE OF SEQUENC CORRESPONDENCE AD ADDRESSEE: STE STREET: 100 NE STREET: 100 NE STREET: 100 NE COMPUTER READABLE MEDIUM TYPE: F COMPUTER: EADABLE MEDIUM TYPE: F COMPUTER: EADABLE MEDIUM TYPE: F COMPUTER: EADABLE MEDIUM TYPE: COMPUTER: EBM OPERATION NUM FILING DATE: 2 CLASSIRICATION NUM FILING DATE: 2 ATTONNEY/AGENT IN NAME: MILLAMN, NAME: MILLAMN, REGISSTRATION NUM RE
228 33321 23321 4444 4444 4444 4444 4444 44		RESULT OS-640-31-6 Sequence Patent N GENERAL APPLIC TITLE TITLE TITLE TITLE COM STA STA STA SCOM APPLIC COM APPLIC APPLIC COM APPLIC APPLIC COM
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241 TIGCIGAACIGIGAGGAGGAAGGCAAITIGAAGACCIIGICAAGGAIAIAACGIIA
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                              GAGAGAAICTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
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  Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/08858197
Patent No. 6297052
GENERAL INFORMATION
APPLICANT: KEHRY, MERLLYN R
APPLICANT: CASTLE, BRIAN E
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   Conservative
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793 TGA 795
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AACAAAGAAGAAAAAAAGAAAACAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAA ATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC **AAGAAAGGATATTATACCATGAAAAGCAACTTGGTAATGCTTGAAAATGGGAAACAGCTG**

CTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180

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ATGAAAATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG
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                                                     Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   97.1%; Score 760.6; DB 4;
98.2%; Pred. No. 1.6e-220;
1ve 0; Mismatches 14;
                                                                                                                                                                                                                         1011.1030000/RAM
                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/234,580,
FILING DATE: 28-APR-1994
ATTORNEY/AGENT INFORMATION:
                                                               CURRENT APPLICATION DATA: US/08/858,197
RILING DATE:
                                        SYSTEM: PC-DOS/MS-DOS'
Patentin Release #1.0,
              : Floppy disk
IBM PC compatible
                                                                                                                                                                                             NAME: MILLMAN, ROBERT A REGISTRATION NUMBER: 36,217
                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                           (202)371-2600
                                                                                                                                                                                                                                                                         TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                      818 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                        nucleic acid
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                                             OPERATING SYSTEM:
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                                                                                                                       CLASSIFICATION:
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US-08-858-197-1
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Matches 769;
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                                           Query, Match
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                                                                                                                 GAGAGAATCTTACTCAAGGCGGCAAATACCCACAGTTCCTCCCAGCTTTGCGAGCAGCAG
                                                                                                                                                                                                                                                                                                                        APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Version 5.1a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: Apple MacIntosh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, Version
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/446,922
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USSN 08/107,353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: PEFKINS, PATRICIA A
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 1003-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Immunex Corporation
STREET: 51 University Street
                                                                                                                                                                                                                                                                                     Sequence 5, Application US/08446922
Patent No. 5716805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: CDNA to mRNA HYPOTHETICAL: NO
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 08-13-93
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NAME/KEY: CDS
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Score 759; DB 1; 1
Pred. No. 4.8e-220;
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Sequence 1, Application US/08249189
Patent No. 5961974
GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
APPLICANT: SPRIGGS, MELLIAM
APPLICANT: SPRIGGS, MELANIE
                                             0;
96.9%;
98.1%;
                                               Conservative
                         Best Local Similarity
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TCTGTTCACTTGGGCGGAGTGTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG 720
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181 GAAGAITITGTAITCATAAAAAGCTAAAGAGAIGCAACAAAGGAGAAGGAICTITATCC 240
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                                                                                                                                                                                                            241 TTGCTGAACTGTGAGGAGATGAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA
                                                                                361 ATTGCAGCACATGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC
                 Sequence 1, Application US/08484624A Patent No. 5962406
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FANSLOW, WILLIAM
SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
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                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintos
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARRE: Microsoft Word for Apple, version 5.1a
                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/249,189
 APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: December 5, 1991
CLASSIETCATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
                                                                                S: IMMUNEX CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 October 25, 1991.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                           December 5, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRANDEDNESS: single
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                                                 NUMBER OF SEQUENCES: 2 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                CITY: SEATTLE
STATE: WASHINGTON
COUNTRY: USA
ZIP: 98101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: cDNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IMMEDIATE SOURCE
CLONE: CD40-L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                   FILING DATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION:
US-08-249-189-1
                                                                                      ADDRESSEE:
                                                                                                  STREET:
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SYĞTEM: Apple Operating System 7.5.5 'NEW Word for Apple Power Macintosh, version 6.0.1 APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
APPLICANT: MCGREW, JEFERX
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
CORRESPONDENCE ADDRESS: E: IMMUNEX CORPORATION 51 UNIVERSITY STREET MEDIUM TYPE: Floppy disk COMPUTER: Apple Macintosh CURRENT APPLICATION DATA COMPUTER READABLE FORM: ADDALL STREET: 51 CITY: SEATTLE CITY: WASHINGTON OPERATING SYSTEM: 98101

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1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60

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FILING DATE: May 24, 1994
CLASSIFICATION: 435
IOR APPLIANT
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APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
CLASSIFICATION: 435
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CORRESPONDENCE ADDRESS:
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TGA 783
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US-08-477-733B-1
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Pred. No. 4.8e-220;
0; Mismatches 15;
US/08/484,624A
                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
CLASSIFICATION:
PRIOR APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
CLASSIFICATION:
                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                            APPLICATION NUMBER: 08/477 FILING DATE: June 07, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               96.9%;
98.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 783 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 96.9
Best Local Similarity 98.1
Matches 768; Conservative
                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear MOLECULE TYPE: CDNA HYPOTHETICAL: NO
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IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                                                                   CLASSIFICATION:
                                                                         FILING DATE: J CLASSIFICATION:
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361 ATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC
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MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: MS Word for Apple Power Macintosh, version 6.0.1
CURRENT Apple.
APPLICATION NUMBER: US/08/477,733B
FILING DATE: June 07, 1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/249,189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/08477733B
Patent No. 5981724
Patent INFORMATION:
APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW WILLIAM
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
APPLICANT: GIBSON, MARYLOU
APPLICANT: MCGREW, JEFFERY
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ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC
                                            601 GAGAGAATCTTACTCAAGGGGGCAAAATACCCACAGTTCCTCCCAGCTTGCGAGCAGCAG
                                                           TCTGTTCACTTGGGCGGAGTGTTTGAATTACAAGCTGGTGCTTCTGTGTTTGTCAACGTG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk COMPUTER: Apple Meditosh OPERATING SYSTEM Maphe Operating System 7.5.5 SOFTWARE: MS Word for Apple Power Macintosh, version 6.0.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40 NUMBER OF SEQUENCES: 26 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/088,913A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: June 07, 1995
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                           FANSLOW, WILLIAM
SPRIGGS, MELANIE
SRINIVASAN, SUBHASHINI
GIBSON, MARYLOU
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December 5, 1991
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                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/09088913A Patent No. 6087329 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             IMMUNEX CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   October 23, 1992
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           51 UNIVERSITY STREET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    MORRIS, ARVIA E. MCGREW, JEFFERY
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FILING DATE: Decemb
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FILING DATE: OCTOD
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TGA 783
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Pred. No. 4.8e-220;
0; Mismatches 15; Indels
                                  CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
                    December 5, 1991
                                                                                                                         REGISTRATION NUMBER: 34,693
REGISTRATION NUMBER: 24,693
REFERENCE/POCKET NUMBER: 286
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 96.9%;
Best Local Similarity 98.1%;
Matches 768; Conservative
                                                                                         CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                   2065870606
                                                                                                                                                                                                                                                                                          linear
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IMMEDIATE SOURCE:
CLONE: CD40-L
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HYPOTHETICAL:
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                                                                                                                                                                                                                                               Length 783
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98.1%; Pred. No. 4.8e-220;
iive 0; Mismatches 15;
LENGTH: 783 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                          Query Match 96.9
Best Local Similarity 98.1
Matches 76%; Conservative
                              single
                                                       MOLECULE TYPE: CDNA
                                          linear
                                                                                                                                                                                    1..783
                                                                                                             ORGANISM: MOUSE
IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                  ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: MOU
                           STRANDEDNESS:
                                                                   HYPOTHETICAL:
                                        TOPOLOGY:
                                                                                                                                                                       NAME/KEY:
                                                                                                                                                                                   ; LOCATION:
US-09-088-913A-1
                                                                                                                                                        FEATURE:
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APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSIOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
ATITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                      SYSTEM: Apple Operating System 7.1 Microsoft Word for Apple, version 5.1a
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19-DEC-1996
18. 424
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
PRIOR APPLICATION NUMBER: 07/805,723
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/788,707
                                                                                                                                                                                                                                                SEE: IMMUNEX CORPORATION: 51 UNIVERSITY STREET SEATTLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 07/783,707
FILING DATE: OCTOBER 25, 1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION WUMBER: 08/484,624
FILING DATE: 07-JUN-1995
APPLICATION NUMBER: 08/249,189
                                      Sequence 1, Application US/08769819
Patent No. 6264951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 26
TELECOMMUNICATION
TELEPHONE: 2065870430
TELEPHONE: 2065870606
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OPERATING SYSTEM: Apple OF
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                                                                                                                                                                                                                                                                                                                                                  ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 783 base pair
TYPE: nucleic acid
                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                  STATE: WASHINGTON
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ORIGINAL SOURCE:
ORGANISM: MOUSE
IMMEDIATE SOURCE:
CLONE: CD40-L
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MOLECULE TYPE: CI
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US-08-769-819-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE:
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RESULT 8
US-08-769-819-1
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CORRESPONDENCE ADDRESS

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ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC 780
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                                                                                        61. AIGAAAAȚITITAIGIAITIACTIACTIACTITICCIIAICACCCAAAIGAITGGAICTGIG 120
                                                                                                   61 ATGAAGATTTTATGTATTACTTACTGTTTTCCTTATCACCCAAAGATGGTTGGATCGTTGT 120
                      0; Gaps
                                          1 ATGATCGAAACATACAACCTACTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
                                                          ACGGTTAAAAGAGAAGGACTCTATTATGTCTACACTCCAAGTCACCTTCTGCTCTAATCGG
Length 783;
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Score 759; DB 4;
Pred. No. 4.8e-220;
                        0; Mismatches
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APPLICANT: SPRIGGS, MELMINE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVE: CVANCON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application US/08770974 Patent No. 6290972
  Query Match 96.9%;
Best Local Similarity 98.1%;
Matches 768; Conservative
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APPLICANT: ARMITAC
APPLICANT: FANSLOI
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TGA 783
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                                                                                                                         CURRENT APPLICATION SYSTEM: Apple Operating System 7.1 SOFTWARE: Microsoft Word for Apple, version 5.1a CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/770,974 FILING DATE: 20-DEC-1996 CLASSIFICATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 96.9%; Score 759; DB 4; I 98.1%; Pred. No. 4.8e-220; Ilve 0; Mismatches 15;
                                                                                                                                                                                                                                                                            AFFLICATION NUMBER: 08/249,189
FILLING DATE: May 24, 1994
                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
5: IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 08/477,733
                                                                                                                                                                                                                                                                                                                                                                                           07/805,723
                                                                                                                                                                                                                                                                                                                                                                                                                                            07/783,707
                                                                                                                                                                                                                                                                                                                                                               FILING DATE: October 23, 1992
PRIOR APPLICATION UNBER:
APPLICATION UNBER: 07/805,72:
FILING DATE: December 5, 1991
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: October 25, 1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 783 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER:
                                                    WASHINGTON
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Best Local Similarity
Matches 768; Conserv
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                                      SEATTLE
                                                                       USA
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US-08-770-974-1
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                                                                                      98101
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                                                                       COUNTRY:
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GAGCCTTCGAGTCAACGCCCATTCATCGTCGCCTCTGGCTGAAGCCCAGCATTGGATCT
                                                                                                    AACAAAGAAGAAAAAAAAAAAAAGAGCTTTGAAATGCAAAGAGGTGATGAGGATCCTCAA
                                                                                                                                                               ATTGCAGCACACGTTGTAAGCGAAGCCAACAGTAATGCAGCATCCGTTCTACAGTGGGCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATER: PATEM: PC-DOS/MS-DOS
SOFTWARE: PATEMIIN Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 5, Application PC/TUS9310034
GENERAL INFORMATION:
APPLICANT: Springs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing
TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/10034
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunex Corporation
51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A
RECISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 1003
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STREET: 51 Univ
CITY: Seattle
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CLASSIFICATION:
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TGA 783
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PCT-US93-10034-5
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61 ATGAAGATTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGT 6120
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4.8e-220;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      96.98;
98.18;
                                                                                                                                                                              CDNA to mRNA
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 5:
EBOUENCE CHARACTERISTICS:
LENGTH: 783 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               768; Conservative
                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 768; Conserv
                                                                                                                                                                                                                                                                    Mouse
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S
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                                                                                                                                                                                                                                                                                                                            ; NAME/KEY: CDS
; LOCATION: 1..7
PCT-US93-10034-5
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                                                                                                                                                                              MOLECULE TYPE:
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                                                                                                                                                   TOPOLOGY:
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                                                                                                                                                                                                                                                                                                               FEATURE
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691 CAAGCTGGTGCTTCTGTGTTTGTCAACGTGAAGCAAGCCAAGTGATCCACAGAGTT 750
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                                                                                                                                                                                 571 GGCCTCTGGCTGAAGCCCAGCATTGGATCTGAGAGAATCTTACTCAAGGCGGCAAATACC
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                                                                                                                                                                                                                           GATAAGGTCGAAGAAGTAAACCTTCATGAAGATTTTGTATTCATAAAAAAGCTAAAG
                                                                                                                                                                   211 AGATGCAACAAAGGAGAAGGATCTTTATCCTTGCTGAACTGTGAGGAGAAGGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: STRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
APPLICANT: MORRELS, ARVIA E.
APPLICANT: MOGREM, JEFERRY
APPLICANT: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 26
                                                  Score 631.4; DB 2;
Pred. No. 2e-181;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            751 GGCTTCTCTTTTGGCTTACTCAAACTCTGA 783
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51 UNIVERSITY STREET
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SPRIGGS, MELANIE
                                                  80.6%;
milarity 99.8%;
Conservative 0
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93..857
                                                      Query Match
Best Local Similarity
Matches 632; Conserv
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US-08-484-624A-22
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   LOCATION:
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             US-08-249-189-22
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    721 ACTGAAGCAAGCCAAGTGATCCACAGAGTTGGCTTCTCATCTTTTGGCTTACTCAAACTC 780
                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Machincoh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                         APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/249,189
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
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                                                                                                                                                                                                                                                                                                                              5: IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                     Sequence 22, Application US/08249189
Patent No. 5961974
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FILING DATE: December 5, 1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 286 FELECOMMUNICATION INFORMATION: 2065870430
                                                                                                                                                                                                                ARMITAGE, RICHARD
FANSLOW, WILLIAM
SPRIGGS, MELANIE
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SEQUENCE CHARACTERISTICS:
LENGTH: 878 base pairs
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15..857
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                                                                                                                                                                                                                                                                                                                                                                                           USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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                                                                     781 TGA 783
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-08-249-189-22
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CACAGTTCCTCCCAGCTTTGCGAGCAGCAGTCTGTTCACTTGGGGCGGAGTGTTTGAATTA 690
                    465 AGTAATGCAGCATCCTTCTACAGTGGGCCAAGAAAGGATATTATACCATGAAAAGCAAC 524
                                                                                                   705 CACAGTTCCTCCCAGCTTTGCGAGCAGTCTGTTCACTTGGGCGGAGTGTTTGAATTA 764
                                                                                                                                                                                                                                                                                                                                                                                                                          AGTAATGCAGCATCCGTTCTACAGTGGGCCAAGAAAGGATATTATATACCATGAAAAGCAAC 450
                                                                                                                                                                                                   585 TACACTCAAGTCACCTTCTGCTCTAATCGGGAGCCTTCGAGTCAACGCCCATTCATCGTC 644
                                                                                                                                                            511 TACACTCAAGTCACCTTCTGCTCTAATCGGGAGCCTTCGAGTCAACGCCCATTCATCGTC
                                                                                                                                                                                                                                             GGCCTCTGGCTGAAGCCCAGCATTGGATCTGAGAGAATCTTACTCAAGGCGGCAAATACC
                                                                                                                                                                                                                                                                                                                                                                                                      TTGGTAATGCTTGAAAATGGGAAACAGCTGACGGTTAAAAAGAGAAGGACTCTATTATGTC
                                                                                                                                                                                COMPUTER: Apple Mcintosh Operating System 7.5.5 OPERATING SYSTEM: Apple Operating System 7.5.5 OPERATING SYSTEM: Apple Operating System 7.5.5 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/477,733B FILING DATE: June 07, 1995 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40 WIMBER OF SEQUENCES: 26 CORRESPONDENCE ADDRESS: ADDRESSEE: IMMUNEX, CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 751 GGCTTCTCTTTTGGCTTACTCAAACTCTGA 783
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T: SPRIGGS, MELANIE
T: SRINIVASAN, SUBHASHINI
T: GIBSON, MARYLOU
T: MORRIS, ARVIA E.
T: MCGREW, JEFFERY
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; Sequence 22, Application US/08477733B
; Patent No. 5981724
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FILING DATE: December 5, 1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/24
FILING DATE: May 24, 1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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PRIOR APPLICATION DATA:
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                                                      OPERATING SYSTEM: Apple Operating System 7.5.5 . SOFTWARE: MS Word for Apple Power Macintosh, version 6.0. CURRENT APPLICATION DATA:
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Pred. No. 2e-181;
0; Mismatches 1;
                                                                                                                   APPLICATION NUMBER: US/08/484,624A
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APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
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APPLICATION NUMBER: 07/783,70
FILING DATE: October 25, 1991
                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: December 5, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
RESISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEPRA: 2065870606
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477
FILING DATE: June 07, 1995
CLASSIFICATION:
                                    COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Op
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 80.6%;
Best Local Similarity 99.8%;
Matches 632; Conservative
                      Floppy disk
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93..857
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15..92
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 0
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COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                             RESULT 14
US-09-088-913A-22
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APPLICANT:
APPLICANT:
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pred. No. 2e-181;
0; Mismatches 1;
                                     ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFREENCE/DOCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAX: 2065870606
       APPLICATION NUMBER: 07/783,70
FILING DATE: October 25, 1991
CLASSIFICATION: 435
                                                                                                                                                                                                                 Murine CD40-L trimer
                                                                                                            INFORMATION FOR SEQ 1D NO: 22: SEQUENCE CHARACTERISTICS: LENGTH: 878 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                   80.6%;
nilarity 99.8%;
Conservative
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                                                                                                                                                                                                                                      NAME/KEY: sig_peptide LOCATION: 15..92
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15..857
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Matches 632; Conserva
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LOCATION:
FEATURE:
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; LOCATION:
US-08-477-733B-22
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SOFTWARE: MS Word for Apple Operating System 7.5.5 SOFTWARE: MS Word for Apple Power MacIntosh, version 6.0.1 APPLICATION DATA: APPLICATION NUMBER: US/09/088 01.0. FILING DATF:
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                                            751 GGCTTCTCATCTTTGGCTTACTCAAACTCTGA 783
                                                                              825 GGCTTCTCATCTTTGGCTTACTCAAACTCTGA 857
                                                                                                                                                                            Sequence 22, Application US/09088913A Patent No. 6087329
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
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GIBSON, MARYLOU
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,72
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,70
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ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 26
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
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CLONE: Murine CD40-L trimer
                                                                                                                                                                                                                                                                                                                                                                                                                                          51 UNIVERSITY STREET
                                                                                                                                                                                                                                                    FANSLOW, WILLIAM
SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                              , ARVIA E.
                                                                                                                                                                                                                                                                                                                                APPLICANT: MORRIS, ARVIA E
APPLICANT: MCGREW, JEFERY
TITLE OF INVENTION: NOVEL
UMMBER OF SEQUENCES: 26
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TYPE: nucleic acid
STRANDEDNESS: single
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APPLICATION NUMBER:
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APPLICATION NUMBER:
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APPLICANT: ARMITA
APPLICANT: FANSLO
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285 AGATGCAACAAAGGAGGAAGGATCTTTATCCTTGCTGAACTGTGAGGAGATGAGAAGGCAA 344
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                                                                                    Length 878;
                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
APPLICANT: ARMITAGE, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                  Score 631.4; DB 3;
Pred. No. 2e-181;
0; Mismatches 1;
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                                                                                  Query Match
Best Local Similarity 99.8%;
Matches 632; Conservative
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93..857
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US-09-088-913A-22
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FEATURE
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Pred. No. 2e-181;
0; Mismatches 1;
                                OPERATING SYSTEM: Apple Operating System 7.1
                                                 Microsoft Word for Apple, version
                                                      19-DEC-1996
                                                                                                                                                                                                    FILING DATE: May 24, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: OCTOBER 23, 1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
                                                                                                                                                     APPLICATION NUMBER: 08/484,624 FILING DATE: 07-JUN-1995 APPLICATION NUMBER: 08/249,189
                                                                                                                                                                                                                                           07/969,703
                                                                                                                                                                                                                                                                                              07/805,723
                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 07/783,707
                                                                                                                                                                                                                                                                                                              FILING DATE: December 5, 1991 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                 RILING DATE: October 25, 1991
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IMMEDIATE SOURCE: , CLONE: Murine CD40-L trimer
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TELECOMMUNICATION INFORMATION
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99.8%;
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
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nucleic acid
                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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CITY: SEATTLE STATE: WASHINGTON

COUNTRY:

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Search completed: May 30, 2002, 02:51:38 Job time: 11602 sec

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Run on:

Searched:

Database

Sequence:

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AX090039 Sequence
215017 H. sapiens m
X67878 H. sapiens m
L07414 Human CD40-
X68550 H. sapiens T
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AF116582 Rattus no
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 AR076926 Sequence
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AF344859 Macaca mu
X96710 H.saplens m
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AR169238 Sequence
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Srinivasan, S. and Spriggs, M.K.
Methods of preparing soluble, oligomeric proteins
Patent: US 5716805-A 3 10-FEB-1998;
Location/Qualifiers
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                                                                                                                                                                    1 ATGATAGAAACATACAGCCA.....TTGGCTTACTCAAACTCTGA 786
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                                                                                  May 30, 2002, 02:48:59 ; Search time 6499.83 Seconds
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Listing first 45 summaries
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Kehry,M. and Castle,B.
Methods for proliferating and differentiating B cells with
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1 (bases 1 to 840)
Armitage, R.J., Fanslow, W.C., Spriggs, M.K., Srinivasan, S.,
Gibson, W.G., Morris, A.E. and McGrew, J.T.
Recombinant soluble CD40 ligand polypeptide and pharmaceutical
composition containing the same
Patent: US 5962406-A 11 05-0CT-1999;
Location/Qualifiers
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Local Similarity 98.2%; Pred. No. 4.7e
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                                                                                                                                                                                                    Length 840;
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Armitage, R. J., Fanslow, W. C. and Spriggs, M.K.
Monoclonal antibodies to CD40 ligand, pharmaceutic
comprising the same and hybridomas producing the Patent: US 5961974-A 11 05-OCT-1999;
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	QY 481 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT 540	QY 541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCTCTGCCTAAAGTCCCCGGGTAGA 600	OY 601 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 660 Db 646 TTCGAGAGAGATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 705	OY 661 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720	OY 721 GTGACTGARCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780	OY 781 CTCTGA 786 111111 Db 826 CTCTGA 831	AR103375	5.1 GI:12814963	NISM	UL-2000;	Arganism="unknown" PASE COUNT 266 a 185 c 175 g 214 t ORIGIN	Query Match 97.2%; Score 763.6; DB 6; Length 840; Best Local Similarity 98.2%; Pred. No. 4.7e-193; Matches 772; Conservative 0; Mismatches 14; Indels 0; Gaps 0;	OY 1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGGGC 60	OY 61 ATGAAGATTTTATGATATTTACTTACTGTTTTTCTTATCACCAGATGATTGGTCAGCA 120	OY 121 CTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGGAATCAATTGAT 180	OY 181 GAAGATTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC 240	OY 241 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300 11111111111111111111111111111111111	OY 301 AACAAAGAGGAGGAGAAAAAAAAAAAAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 360
AATACCCACAGTTCGGCCAAACCTTGCGGGCAA 660	TICGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGCCAA 705 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 705		GTGATGATCGAGCCAAGCCAAGCCATGCCTTCAGGTTTGGTTG		840 bp DNA linear PAT 01-SEP-2000 US 5981724.		Unknown. Unclassified. 1 (bases 1 to 840) 1 (bases 1, Fanslow, W.C., Spriggs, M.K., Srinivasan, S.,	a cytokine that binds CD40 09-NOV-1999; iers	own. 5.g 214 t	Score 763.6; DB 6; Length 840; Pred. No. 4.7e-193; 0; Mismatches 14; Indels 0; Gaps 0;	1 ATGATAGAAACATACAGCCAACTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGGGC 60	ATGAAGATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCACA 120 	CTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGGAAG	GAAGATTTGTATTCATGAAAACGATACAGAGATGCAACACAGAGAAAGATCCTTATCC 240 	TTACTGAACTGTGAGGAGTTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300 	aacaaagaggagacgaagaaagaaacagctttgaaatgcaaaaaggtgatcagaatct 360 	CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 420	421 GCTGAAAAAGGATACTACACCATGAGCAACTTGGTAACCCTGGAAAATGGGAAACAG 480
TTCGAGAATCTTACTCAGAGCTGCAA	CGAGAGATCTTACTCAGAGCTGCAA	ATCCATTCACTTGGGAGGAGTALTTC	GACTGATCCAAGCCAAGTGAGCCATG 	781.CTCTGA 786 826.CTCTGA 831	from patent	AR085419.1 GI:10012188	Unknown. Unclassified. 1 (bases 1 to 840) Armitage, R.J., Fanslow, W.	LE DNA encoding CD40 ligand, as RNAL Patent: US 5981724-A 11 09-N RES LOCATION/Qualifiers Source 1. 840	/organism="unknown" 266 a 185 c 175 g	Query Match Q7.2%; Scc Best Local Similarity 98.2%; Pre Matches 772; Conservative 0;	TGATAGAAACATACAGCCAACCTTC(ngaagatitiatctatitacttac 	CTTTTGCTGTGTATCTTCATAGAAG	SAAGATTTGTATTCATGAAAACGAT 	ttactgaactgtgaggagattaaaag 	aacaaagaggaggcgaagaagaaaa 	CAAATTGCGGCACATGTCATAAGTGP 	GCTGAAAAAGGATACTACACCATGAG

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                                         TTACTGAACTGTGAGGAGAGTTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA
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Sequence 11 from patent US 6290972.
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Spriggs,M.K., Armitage,R.J., Fanslow,W.C. III and Widmer,M.B.
Treatment of individuals exhibiting defective CD40L
Patent: US 6106832-A 7 22-AUG-2000;
Location/Qualifiers
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                                                                                                                 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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Sequence 7 from patent US 6106832.
AR106246.1 GI:12820776
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PAT 17-DEC-2001 Gaps by administering CD40 9 and 1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC 1 (bases 1 to 840) Armitage.R.J., Fanslow,W.C., Spriggs,M.K., Srinivasan,S. Gibson,M.G. ö linear Length Indels Score 763.6; DB 6; Pred. No. 4.7e-193; 0; Mismatches 14; Method of augmenting a vaccine response

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Spriggs,M.K., Armitage,R.J. and Fanslow,W.C.
Detection of mutations in a CD40 ligand gene
Patent: US 556531-A 7 15-OCT-1996;
                                   DNA
                                 840 bp
US 5565321
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                     Aruffo, A. Hollenbaugh, D. and Ledbetter, J. A. Soluble and its use in B cell stimulation Patent: US 5540926-A 1 30-JUL-1996; Location/Qualifiers
                                                                                                                  DB 6;
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Pred. No. 4.7e-193;
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k; Pred. No. 4.7e-193;
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Catarrhini;
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/db_xref="taxon:9606"
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AX090039
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Mammala; Eutheria; Primates;
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Ahuja,S.U. and Bonewald,L.U.
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                                   GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780
                                                 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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Pred. No. 4.7e-193;
0; Mismatches 14;
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Armitage, R.J., Fanslow, W.C. III, Longo, D.
Method of preventing or treating disease
method objastic cells expressing CD40
patent: US 5674492-A I 07-OCT-1997;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Hollenbaugh,D., Grosmaire,L.S., Kullas,C.D., Chalupny,N.J.,
Braesch-Andersen,S., Noelle,R.J., Stamenkovic,I., Ledbetter,J.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              and Aruffo, A.
The human T cell antigen gp39, a member of the TNF gene family, a ligand for the CD40 receptor: expression of a soluble form of gp39 with B cell co-stimulatory activity
EMBO J. 11 (12), 4313-4321 (1992)
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Submitted (21-SEP-1992) Diane L Hollenbaugh,
PRI, Seattle, WA, 98121
The original submission [1] reported 934bp.
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mRNA for glycoprotein 39
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/protein_id="CAA78737.1"
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/db_xref="taxon:9606"
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glycoprotein 39.
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Hollenbaugh, D.L.
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/db_xref="SWISS-PROT:P29965"
/translation="MIETYNOTSPRSAATGLPISMKIFMYLLTVFLITQMIGSALFAV
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EETKKROKELSEMGKGDONPQIAAHVISBASSKTTSVLQWAEKGYTMSNNLYTLENGKQ
LIVKROGLYYIYAQYTKSNREASSQAPFIASLCLKSPORRERILLRAANTHSSAKPC
GQOSIHLGGVFELQPGASVFVNVTDPSQVSHGTGFTSFGLLKL"

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Pred. No. 4.7e-193;
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/db_xref="GI:38484"
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98.2%;
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YLHRRLDKIEDERNIHEDFYPMKTIORCNTGERSLSLLINCEEIKSOFEGFVKDIMLNK
EETKKENSFEMOKGDONPOIAAHVISEASSKTTSVLQWAEKGYYTMSNNLYTLENGKO
LTVKROGLYXIYAQYTFCSNREASSOAPFIASLCLKSPGRFERILLRAANTHSSAKPC
     PRI 06-JUN-1997
                                                                                                                                                                                                     Spriggs, M.K., Armitage, R.J., Strockbine, L., Clifford, K.N., Macduff, B.M., Sato, T.A., Maliszewski, C.R. and Fanslow, W.C. Recombinant human CD40 ligand stimulates B cell proliferation and immunoglobulin E secretion
                                                                                                   Euteleostomi;
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                                                                                                                                                            Direct Submission
Submitted (28-Jul-1992) M. Spriggs, Immunex Res & Development
Submitted (28-Jul-1992) M. Spriggs, Immunex Res & Development
Corporation, 51 University Street, Seattle WA 98101, USA
2 (bases 1 to 1803)
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1803)
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Pred. No. 4.5e-193;
); Mismatches 14;
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/db_xref="taxon:9606"
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18 46.8 6.0 1043 12 CNS0145P AL164525 AL164525	ALIGNMENTS	RESULT 1 BF599437 492 bp mRNA linear EST 25-APR-2001 LOCUS BF599437 DEFINITION 263218 MARC 3BOV Bos taurus cDNA 5', mRNA sequence. ACCESSION BF599437 VERRION BF599437.1 GI:11695919 SOURCE COW. SOURCE COW. ORGANISM BOS taurus	ء آن س <u>ـ</u>	Casas, E., Waray, J.E., White, J., Cho, J., Fahrenkrug, S.C., G.L., Heaton, M.P., Laegreid, W.W., Rohrer, G.A., Chitko-Pertea, G., Holt, I., Karamycheva, S., Liang, F., Quackenb Keele, J.W.	TITLE Sequence evaluation of four pooled-tissue normalized bovine CDNA libraries and construction of a gene index for cattle JUDURNAL Genome Res. 11 (4), 626-630 (2001) MEDLINE 21180013 COMMENT CONTACT: Smith TPL USDA, ARS, US Meat Animal Research Center PO Box 166, Clay Center, NE 68933-0166, USA Tel: 402 762 4360 Fax: 402 762 4360 Email: smith@email.marc.usda.gov Single pass sequencing. Bases called and alt_trimmed with phred v0.980904.e. Vector identified by cross match with the -minscore 18	b."
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4.5 Compugen Ltd. Trch time 5112.58 Seconds (without alignments) 2075.001 Million cell updates/secTTGGCTTACTCAAACTCTGA 786 sidues ters: 27472414	*			cted by chance to have a soft the result being printed, score distribution.	Description BF59437 263218 MA AW486605 75217 MAR AJ982044 par.pk007 AZ166561 SP_0088_B AL060767 Drosophil BF820152 MR1-RT002 AV714552 AV714252	AV724813 AV724813 BI711310 160251255 AL534423 AL534423 BG715238 602676117 BI601432 603249012 BF577761 602077072 BG616023 602643466 BG715241 602676120 AA312300 EST182997
GenCore version 4. pyright (c) 1993 - 2000 Coo search, using sw model 30, 2002, 01:00:10 ; Searc 08-982-272-6 TGATAGAAACATACAGCCA NTITY_NUC op 10.0 , Gapext 1.0 36207 seqs, 6748477542 resi satisfying chosen paramete h: 0 h: 2000000000 imum Match 0% imum Match 100% ting first 45 summaries	EST:		13: em_gss_hum:* 14: em_gss_lnv:* 15: em_gss_phn:* 16: em_gss_vrt:*	he number of results prediction or equal to the score by analysis of the total	Ch Length DB 492 10 12 19 638 9 14 1027 12 997 12 12 102 12 102 12 102 12 102 102 102	6.0 578 9 AV724813 6.0 583 10 BF73130 6.0 586 10 BF731160 6.0 797 9 AL53423 6.0 845 10 BG71233 6.0 845 10 BF671233 6.0 873 10 BF673761 6.0 874 10 BG715241 6.0 553 9 AA312300
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Boyidae; Boxinae; Bos.

I (bases I to 399)

Smith, T.P.L., Grosse, W.M., Freking, B.A., Roberts, A.J., Stone, R.T., Casas, E., Wray, J.E., White, J., Cho, J., Fahrenkrug, S.C., Bennett, G.L., Heaton, M.P., Laegreid, W.W., Rohrer, G.A., Chitko-McKown, Cif. Pertea, G., Holt, I., Karamycheva, S., Liang, F., Quackenbush, J. and Keele, J.W.
                                   /note="Vector: pcwv SPORT6; Site_1: Xbal; Site_2: Xhoi; //note="Vector: pcwv SPORT6; Site_1: Xbal; Site_2: Xhoi; // Library made from pooled tissue from marrow, alveolar macrophage, ovary, fetal semitendonosus muscle, and fetal longissimus muscle = 12 t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence evaluation of four pooled-tissue normalized bovine cDNA libraries and construction of a gene index for cattle Genome Res. 11 (4), 626-630 (2001)
                                                                                                                                                                                                                                                                                                                                                                       361 CAAATIGGGGACAIGICATAAGIGAGGCAGCAGCAGAAAAACAACAICIGIGITACAGIGG 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATGAAGATTTTTATGTATTTACTTACTGTTTTTTTTTATCACCCAGATGATTGGGTCAGCA 120
                                                                                                                                                                                                                                                                            CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
                                                                                                                                                                                                                                                                                                                                   TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
                                                                                                                                                                                                                                                                                                                                                                                                                                              301 AACAAAGAGGAGGAGAAAAAAAAAAAAGCTTTGAAATGCAAAAAAGGTGATCAGAATCCT 360
                                                                                                                                                                                                                    46 AIGATCGAACATCAACCTTCTCCCGCTCGTGGCCACTGGACCACTGTCAGT 105
                                                                                                                                                                            Gaps
                                                                                                                                                                                                     1 ATGATAGAAACATACAGCCAAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC 60
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0
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5', mRNA sequence:
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USDA, ARS, US Meat Animal Research Center
PO Box 166, Clay Center, NE 68933-0166, USA
                                                                                                                                                  Score 359; DB 10;
Pred. No. 2.3e-71;
0; Mismatches 55;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Craniata;
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75217 MARC 1BOV BOS taurus CDNA
/clone_lib="MARC 3BOV"
/tissue_type="pooled"
/lab_host="DH10B"
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                                                                                                                                                   Query Match 45.7%;
Best-Local Similarity 87.7%;
Matches 392; Conservative
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A1982044 638 bp mRNA linear EST 07-MAY-2001 pat.pk0072.c9.f chicken activated T cell cDNA Gallus gallus cDNA clone pat.pk0072.c9.f 5' similar to CD40 ligand, mRNA sequence. A1982044
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
Phasianinae; Gallus.
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Tirunagaru,V.G., Sofer,L., Cui,J. and Burnside,J.
An expressed sequence tag database of T-cell-enriched activated chicken splenocytes: sequence analysis of 5251 clones Genomics 66 (2), 144-151 (2000)
                                                                  -minscore
                                                                                                                                                                                                                                                                                                                               /note="Vector: pcMv SPORT6; Site_1: XbaI; Site_2: XhoI; Library made from pooled tissue from lymph node, ovary, fat, hypothalamus, and pituitary."
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with the -mins
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 398;
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Pred. No. 2.9e-54;
0; Mismatches 39;
                                                    called and
                                                      pass sequencing. Bases call 04.e. Vector identified by
                              Email: smith@email.marc.usda.gov
Single pass sequencing. Bases ca.
v0.980904.e. vector identified by
and -minmatch 12 options.
                                                                                                                                                                                                                                                             /db_xref="taxon:9913"
/clone_lib="MARC 1BOV"
/tissue_type="pooled"
                                                                                                                                             BACKWARD: GTTTTCCCAGTCACGACG Plate: 34 row: F column: 23 seq primer: ATTTAGGTGACACTATAG
                                                                                                                                                                                                                                            /organism="Bos taurus
                                                                                                               PCR PRIMERS
FORWARD: AGGAAACAGCTATGACCAT
                                                                                                                                                                                                        Location/Qualiflers
                                                                                                                                                                                                                                                                                                                     /lab_host="DH10B"
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ilarity 88.7%;
Conservative
Tel: 402 762 4366
Fax: 402 762 4390
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Matches 307; Conserv
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Pasadena California 91125, USA
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,G.A., Ettensohn,C.A., Lehrach,H., Britten,R.J, Davidson,E.H. and
Hood,L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biondi, T.R.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       116 GAAGGGACTTGACACGCACAGCACCTCCTCTGTGAGCTCCAGTCCATCCGGGA 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       558 AGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGATTCGAGAATCTTACT 617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          57 -GGGCCATTCACCCTCTATATTTATTGTACCTCCCCATGGAAGAGGACCGGCTCCTGAT 115
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              498 AGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAATCGGGAAGCTTCGAGTCA 557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 AGGGCTCTACTACATCTCACAAGTCAGCTTCTGCACCAAGGCGGCGGCTTCG----- 56
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              518 CAGAGCTGCAAATACCCACAGTTCCGCCAAACCT---TGCGGGCAACAATCCATTCACTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Proc. Natl. Acad. Sci. U. S. A. 97 (17), 9514-9518 (2000)
20402266
                                                                                                                                                                                                                                                                                                                                                                                                                       Length 638;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       236 AGCAGTGAACGTCAACCTGCCAACATATGGCATGTTCAAGCTGT 285
                                                                                                                                                                                                                                                                                     /cell_type="Con A-activated splenic T cell"
/lab_host="E.coli TOP10 F'"
/note="Vector: pcDNA3"
171 c 151 g 180 t 4 others
                                                                                                                                                                     1.638
/organism="Gallus gallus"
/orb_xref="taxon:9031"
/clone="pat.pk0072.c9.f"
/clone=lpb="chicken activated T cell cDNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 735 CCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAACTCT 784
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Cameron, R.A., Mahairas, G., Rast, J.P., Martinez, P.,
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Eukaryota; Metazoa; Echinodermata; Eleutherozoa;
Echinoidea; Euechinoidea; Echinacea; Echinoida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Cameron, RA, Davidson, EH, Hood, L
               Molecular Endocrinology
University of Delaware
40 Townsend Hall, Newark, DE 19717, USA
Tel: 302 831-1345
Fax: 302-831-3411
Email: joan@UDel.Edu, www.chickest.udel.edu
Seg primer: T7
                                                                                                                                                                                                                                                                                                                                                                                                                     Score 69.6; DB 9;
Pred. No. 1.1e-05;
0; Mismatches 114;
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California Institute of Technology
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                                                                                                                                                     Location/Qualifiers
Contact: Joan Burnside
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997 bp DNA linear GSS 03-JUN-1999 Drosophila melanogaster genome survey sequence TFT3 end of BAC #BACR12K22 of RPCI-98 library from Drosophila melanogaster (fruit AL060767
                                                                                                                                                                                                                                 /organism="Strongylocentrotus purpuratus"
/db_xref='taxon:7668"
/db_xref='taxon:7668"
/clone="Plate=88 Col=3 Row=F"
/clone_lipl="Strongylocentrotus purpuratus, purple sea
urchin, sperm genomic BAC library"
//note="Organ: sperm; Vector: BACe3.6; BAC Clones in E-Coli
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Determination of this BAC-end sequence was carried out as part of a collaboration with the Berkeley Drosophila Genome Project (BDGP).
The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further information please see http://www.fruitfly.org The BDGP Drosophila melanogaster BAC ilbrary was prepared by Kazutoyo Osogawa and haron Mammoser in Pieter de Jong's laboratory in the Department of
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BP 191 91006 EVRY cedex - FRANCE (E-mail : seqref@genoscope.cns.fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             201 AACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCCTTACTGAACTGTGAGGAGAT 260
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Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
1 (bases 1 to 997)
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Pred. No. 0.27;
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Eukaryota; Metazoa; Arthropoda;
                                                                                                                                                            High quality sequence stop: 1
Location/Qualiflers
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                                                 Email: acameron@caltech.edu
Plate: 88 row: F column:
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Tel: (626) 395-8421
Fax: (626) 793-3047
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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Class: BAC ends
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Matches 142; Conserv
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Unpublished (2000)
Contact: Zeguang H
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  EST 13-JAN-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20152
-R70025-171100-003-C06 RT0025 Homo sapiens CDNA, mRNA sequence-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Euteleostomi;
   Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPCI-98 and was constructed by partial scorl digestion of Drosophila DNA provided by the BDGP from the isogenic strain y2; cn bw sp, the same strain used for the BDGP's Pl and EST libraries. A more detailed description of the library and how to order individual BAC clones, the entire library filters for hybridization from the BACPAC Resource Center can be found at http://bacpac.med.buffalo.edu/drosophila_bac.htm.
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Primates, Catarrhini, Hominidae, Homo.
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                                                                                                                                                                                                                                                                                                                  Length 997;
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Conservative 143; Mismatches
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Pred. No. 0.
                                                                                                                                                                      /organism="Drosophila
/db_xref="taxon:7227"
/clone_lib="RPCI-98"
/clone="BACR12K22"
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Mammalia; Eutheria;
1 (bases 1 to 396)
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59; Conserv
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clone DCBADF07.5', mRNA sequence.
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                                                                                                                                                                         Email: asimpson@ludwig.org.br
This sequence was derived from the FAPESP/LICR Human Cancer Genome
This sequence was derived from the FAPESP/LICR Human Cancer Genome
This entry can be seen in the following URL
(http://www.ludwig.org.br/scripts/gethtml2.pl?tl=MR1&t2=MR1-RT0025-
171100-003-c06&t3=2000-11-17&t4=1)
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                                                                    01509-010, Sao Paulo-SP,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           146 CICAAAIACGIGGACACCAAIIIGCAGAAACAGCGAGAACIICAAAAAGCCAAICAGGAG
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0; Mismatches 118;
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                           Laboratory of Cancer Genetics
Ludwig Institute for Cancer Research
Rua Prof. Antonio Prudente 109, 4 andar,
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/db_xref="taxon:9606"
/clone_lib="RT0025"
/dev_stage="Adult"
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High quality sequence start: 8
High quality sequence stop: 396.
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Mammalia; Eutheria; Primates;
1 (bases 1 to 486)
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Contact: Simpson A.J.G.
Laboratory of Cancer Genetics
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                                                                                                                          Tel: +55-11-2704922
                                                                                                                                                           Fax: +55-11-2707001
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1 (bases 1 to 578)

Gu,Y., Cano,Y., Song,H., Huang,Q., Yang,Y., Gao,G., Xiao,H., Xu,X., Li.N., Qian,B., Liu,F., Qu,J., Gao,X., Cheng,Z., Xu,Z., Zeng,L., Xu,S., Gu,W., Tu,Y., Jia,J., Fu,G., Ren,S., Zhong,M., Lu,G., Hu,R., Chen,J., Chen,Z. and Han,Z.

Homo sapiens cDNA HTB clones
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AV724813 578 bp mRNA linear EST 16-OCT-2000 AV724813 HTB Homo sapiens CDNA clone HTBAKC04 5', mRNA sequence.
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Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai
201203, P. R. China
Tel: 86-21-50801919(ex.45)
Fax: 86-21-50801919(ex.45)
Email: hang@ehgqc.sh.cn
This clone is available at CHGC in Shanghai.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Contact: Zeguang Han
Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai
201203, P. R. China
                                                                                                                                                                                                                                                                                     Site_1: sfiIA; Site_2: sfiIB"
t
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                                                                                                                                                                                                                                                                                                                                                                             Length 486;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Email: hanzg@chgc.sh.cn
This clone is available at CHGC in Shanghai.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                         6.0%; Score 47.2; DB 9; 50.0%; Pred. No. 1.3;
                                                                                                                                                                                                                                 cells"
                                                                                                                                                                                                                                                                                     /note="Vector: pTriplEx2;
87 c 124 g 93
                                                                                                                                                      /organism="Homo sapiens"
/db_xref="taxon:9606"
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/db_xref="taxon:9606"
/clone="HTBAKC04"
                                                                                                                                                                                                             /clone_lib="DCB"
/cell_type="dendritic
                                                                                                                                                                                                                                                /dev_stage="mature"
/lab_host="BM25.8"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              201203, P. R. China
Tel: 86-21-50801919(ex.45)
                                                                                                                                                                                              /clone="DCBADF07"
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Matches 118; Conservative
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ORIGIN
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JOURNAL
COMMENT
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AUTHORS
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KEYWORDS
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HEVI3130 583 bp mRNA linear EST 19-SEP-2001 1e02f03.yl HR85 islet Homo sapiens cDNA 5' similar to TR:095899 DINKNOWN; mRNA sequence.
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Melton, D., Brown, J., Kenty, G., Permutt, A., Lee, C., Kaestner, K.,
Lemishka, I., Scearce, M., Barestelli, J., Gradwohl, G., Clifton, S.,
Hiller, L., Marra, M., Pape, D., Wylie, T., Martin, J., Blistain, A.,
Schmitt, A., Theising, B., Ritter, E., Ronko, I., Bennett, J., Cardenas, M., Gibbons, M., McCann, R., Cole, R., Tsagareishvili, R., Williams, T.,
Endocrine Pancreas Consortium
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Harvard University, Howard Hughes Medical Institute
Dept of Molecular and Cellular Biology, 7 Divinity Ave, Cambridge,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fax: 617-495-8557
Email: dmelton@biohp.harvard.edu
Email: dmelton@biohp.harvard.ed by
Email: dmelton@biohp.harvard.ed by
Email: dmelton@biohp.harvard.ed by
Washington University Genome Sequencing Center For information on obtaining a clone please contact: Dr. Hiroshi Inoue
(hinoue@im.wustl.edu)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Euteleostomi;
                                                                                         /note-"Vector: pBluescript sk(-); Site_1: EcoRI; Site_2:
                                                                                                                                                                                                                                                                                                                                                                                              238 TCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   298 TTAAACAAAGAGGAGGACGAAGAAAGAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAT 357
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               148 CAGTATGCAGAGGGGAAGATGATAGGGGCTGCCCCAGGAAAGAATACATCTGGTCTGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            108 CAGAAAAATGTTGAAGTGAAAAGGAAAAGAACAAAACACCTGGAAATGG 263
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae;
                                                                                                                                                                                                                                                                                                                                   Indels
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//db_xref="taxon:9806"
//clone_lib=HR85 islet"
//tissue_type="Purified pancreatic islet"
                                                                                                                                                                                                                                                                                                                                0; Mismatches 118;
                                                                                                                                                                                                                                                                         DB 9;
/tissue_type="Hypothalamus"
/dev_stage="Adult"
/lab_host="SOLR"
                                                                                                                                                                                                                                                                     Score 47.2;
Pred. No. 1.
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High quality sequence stop: 471.
Location/Qualifiers
                                                                                                                                                      б
                                                                                                                                                   148
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Other_ESTs: ie02f03.x1
                                                                                                                                                                                                                                                                  6.0%;
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Matches 118; Conservative
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                                                                                                                           XhoI
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BF791160 586 bp mRNA linear EST 12-JAN-2001 602251255F1 NIH_MGC_81 Homo sapiens cDNA clone IMAGE:4338636 5' pr
             SK(-); Site_1:
                                Not1: Site_2: XhoI; cDNA made by oligo-dT priming.
Size-selected on agarose gel. Average insert size -1kb. 5'
XhoI site was destroyed after directional clouing.
Amplified once. Contact information: Hiroshi Inoue, MD,
Metabolism Div. (Alan Permutt Lab), Washington University
School of Medicine, Box 8127, 660 South Euclid Ave., St.
Louis, MO 63110, Eax: 314-747-2692."

114-36-1916, Fax: 314-747-2692."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /clone_lib="NIH_MGC_81"
/lab_host="DH10B (T1 phage-resistant)"
/note="Crgan: muscle (skeletal), Vector: pDNR-LIB
(clontech); site_1: sfil (gccgcctcggcc); Site_2: Sfil(gccattatggcc); 5, and 3' adaptors were used in cloning as follows: 5' adaptor sequence: 5'-CACGCCCATTATGGCC-3'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              164 CICAAAIACGIGGACACCAAIIIGCAGAAACAGCGAGAACIICAAAAAGCCAAICAGGAG 223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCTCAAATIGCGGCACAIGICAIAAGIGAGGCCAGCAGIAAAACAACAICIGIGIIACAG 417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 238 TCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297
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National Institutes.of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Tissue Procurement: CLONETECH Laboratories, Inc.
CDNA Library Preparation: CLONETECH Laboratories, Inc.
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  418 TGGGCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGG 473
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Bukaryota, Metazoa; Chordata; Craniata; Vertebrata; Butele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
1 (bases 1 to 586)
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                                                                                                                                                                                                                                                                                                                                               Length 583;
                  'note="Organ: Pancreas; Vector: pBluescript
                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         found through the I.M.A.G.E. Consortium/LLNL at:
                                                                                                                                                                                                                                                                                                                                          Score 47.2; DB 10;
Pred. No. 1.3;
0; Mismatches 118;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Plate: LLCM1214 row: n column: 13
High quality sequence stop: 582.
Location/Qualifiers
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'lab_host="DH10B"
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                                                                                                                                                                                                                                                                                                                                               Query Match 6.0%;
Best Local Similarity 50.0%;
Matches 118; Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      mRNA sequence.
BF791160
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BF791160
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ORIGIN
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TITLE
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COMMENT
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KEYWORDS
SOURCE
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Technologies. Contact: Feng Liang Life Technologies of the division of Invitrogen 9800 Medical Center Drive Rockville, Maryland 20850, USA Fax: (1) 301 610 8371 Email: filang@lifetech.com URL: http://dullength.invitrogen com."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note="Organ: Fetal brain; Vector: pCMVSPORT 6; 1st strand cDNA was primed with a NotI-Oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   /clone_lib="LTI_FL013_FBrn1"
/dev_stage="pooled tissue from post conception fetuses (20 week, 24 week and 26 week)"
/lab_host="DH10B"
                          C, or G and N = A, C, G, or T). Average insert size 1.55 kb (range 1.0-4.0 kb). 15/15 colonies contained inserts by PCR. This library was enriched for full-length clones and was constructed by Clontech Laboratories (Palo, Alto, CA). 163 g 118 t
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                                                                                                                                                                                                                                                                                                                                                                             298 TTAAACAAAGGGGGGGGGAGGAAGAAGAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAT 357
             -ATTCTAGAGGCCGAGGCGCCGACATG-dT(30)BN-3' (where B
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BP 191 91006 EVRY cedex - France
Email: segref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         418 TGGGCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGG 473
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                                                                                                                                                                                                                 Length 586;
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                                                                                                                                                                                                                 Score 47.2; DB 10;
Pred. No. 1.3;
0; Mismatches 118;
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/db_xref="taxon:9606"
and 3' adaptor sequence:
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Mammalia; Eutheria;
1 (bases 1 to 797)
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                                                                                                                                                                                                                                             Best Local Similarity 50.08
Matches 118; Conservative.
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Matches 151

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Query Match

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B1601432 845 bp mRNA linear EST 07-SEP-2001 603249012F1 NIH_MGC_96 Homo sapiens cDNA clone IMAGE:5300787 5',
                                                                       pBluescript (8+1): Site_1: BamHI; Site_2: Sall XhoI (gtcgs ); Oligo-dT primed using primer 5'-TTTTTTTTTTTTTTTVN-3', size-selected for average insert size 2.3 kb and normalized to TROT 5. This is a primary library enriched for full-length clones and constructed using the Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. 1 (bases 1 to 845.nci.nih, gov/. NIH-WGC http://mgc.nci.nih,gov/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D.

Email: cgapbs-remail.nih.gov

Tissue Procurement: Miklos Palkovits, M.D., Ph.D.

cDNA Library Preparation: Michael J. Brownstein (NHGRI), Shira

Toshiyuki and Plero Carninci (RIKEN)

Clone distribution: MGC clone distribution information can be found through the IN.M.G.E. Consortium/LLNL at:

http://mage.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      238 TCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           298 TTAAACAAAGAGGAGGAGGAAGAAAGAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAT 357
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/tissue_type="hypothalamus"
/lab_host="DHlob"
/note="Organ: Drain; Vector: pBluescriptR (modified
                                                       /note="Organ: brain; Vector: pBluescriptR (modified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            237 CTCAAATACGTGGACACCAATTTGCAGAAACAGCGAGAACTTCAAAAAGCCAATCAGGAG
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/tissue_type="hypothalamus".
/lab_host="DH10B"
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Pred. No. 1.3;
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Location/Qualifiers
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/clone="IMAGE:5300787"
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Matches 118; Conserv
                                                                                                                                                                                                                                                                                      284
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NIH-WGC http://mgc.ncl.nih.gov/.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D.
Email: cgapbs remail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
CDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            387 GCCCAGCAGTAAAACAACATCTGTGTTACAGTGGGCTGAAAAAGGATACTACACCATGAG 446
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Clone distribution: MGC clone distribution information can be
                                                                                                                                                          GACAAGATAGAAGATGAAAGGAATCTTCATGAAGATTTTGTATTCAŢGAAAACGATACAG 210
                                                                                                                                                                                                                                                             AGATGCAACACAGGAGAAAGATCCTTATCCTTACTGAACTGTGAGGAGATTAAAAGCCCAG 270
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Plate: LLAM10686 row: b column: 05
High quality sequence stop: 830.
                                                       DB 9;
                                                                                                        Mismatches
                                                       6.0%; Score 47.2;
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/db_xref="taxon:9606"
/clone="IMAGE:4798372"
/clone_lib="NIH_MGC_96"
                                                                                  Pred. No.
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                                                                                                        Conservative
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BG715238
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RESULT 12 BG715238 LOCUS

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SOURCE ORGANISM

ACCESSION VERSION KEYWORDS REFERENCE AUTHORS TITLE JOURNAL

COMMENT

FEATURES

Shiraki

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602077072F2 NIH_MGC_62 Homo sapiens cDNA clone IMAGE:4251880 5', MRNA seminance
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/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: skin; Vector: pDNR-LIB (Clontech); Site_1:
Sfil (ggccqctctgqcc); Site_2: Sfil (ggccattatggcc);
Double-stranded cDNA was prepared from cell line RNA. 5'
and 3' adaptors were used in cloning as follows: 5'
adaptor sequence: 5'-CACGGCCATTATGGCC-3' and 3' adaptor
Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
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National Institutes of Health, Mammalian Gene Collection (MGC)
Unpublished (1999)
Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                              ICCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297
                                                                                                                                                                                                                                                                                                                                                                                                                                       329 CAGTATGCAGAGGGGAAGATGAGAGGGGCTGCCCCAGGAAGAAGAAGACATCTGGTCTGCAA 388
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                   DB 10;
                                                                                                                                                                                                                                                                                         0; Mismatches 118;
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plate: LLCMINO'S row: o column: 17
High quality sequence stop: 590.
Location/Qualifiers
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Pred. No. 1.3
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Tissue Procurement: ATCC/DCTD/DTP
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/clone_lib="NIH_MGC_62"
                                                                                                                                                                   243 g
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Best Local Simi
Matches 118;
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TITLE
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ORIGIN
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BF573761
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/lab_host-"DH10B (T1 phage-resistant)"
/note="Organ: testis; Vector: pDNR-LIB (Clontech); Site_1:
SfiI (ggccgcctcggcc); Site_2: SfiI (ggccattatggcc);
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        BG616023 873 bp mRNA linear EST 18-APR-2001
602643466F1 NIH_MGC_61 Homo sapiens CDNA clone IMAGE:4774241 5',
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Double-stranded CDNA was prepared from cell line RNA. 5' and 3' adaptors were used in cloning as follows: 5' and 3' adaptor sequence: 5'-CARGGCCARTAGGCCA' and 3' adaptor sequence: 5'-ATTCTAGAGGCGGGGGGGGGGGGATG-dT(30)BN-3' (where B = A, C, or G and N = A, C, G, or T). Average
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Bukaryota; Metazoa; Chordata; Craniata: Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 873)
              (where B = A, C, or G and N = A, C, G, or T). Average insert size 1.75 kb (range 0.9-4.0 kb). 15/15 colonies contained inserts by PCR. This library was enriched for Lull-length clones and was constructed by Clontech Laboratories (Palo Alto, CA). 1 others
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Clone distribution: MCC clone distribution information can be
found through the I.M.A.G.E. Consortium/LLNL at:
http://image.llnl.gov
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NIH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
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                                                                                                                                                                                                                                                                                                                                                                5'-ATTCTAGAGGCCGAGGCGGCCGACATG-dT(30)BN-3'
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                                                                                                                                                                                                                          DB 10; Length 861;
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/clone_11b="NIH_MGC_61"
/tissue_type="embryonal carcinoma"
                                                                                                                                                                                                                            Score 47.2; DB 10;
Pred. No. 1.3;
0; Mismatches 118;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Contact: Robert Strausberg, Ph.D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /db_xref="taxon:9606"
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Tissue Procurement: ATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mRNA sequence.
BG616023
BG616023.1 GI:13667394
                                                                                                                                                                                                                                 Query Match 6.0%;
Best Local Similarity 50.0%;
Matches 118; Conservative
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insert size 1.75 kb (range 0.9-4.0 kb). 15/15 colonies contained inserts by PCR. This library was enriched for full-length clones and was constructed by Clontech Laboratories (Palo Alto, CA). Note: this is a NIH_MGC Library."
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; 0 Gaps ; 0 Length 873; Indels Score 47.2; DB 10; Pred. No. 1.3; 0; Mismatches 118; Query Match 6.0%; Best Local Similarity 50.0%; Matches 118; Conservative (

238 TCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297 셤

298 TTAAACAAAGAGGAGACGAAGAAAAAAAAGCTTTGAAATGCAAAAAGGTGATCAGAAT 357 g

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418 TGGGCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGG 473 ò

Search completed: May 30, 2002, 01:00:13 Job time: 22223 sec

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AAV61063

DNA encoding novel Genomic sequence o Human CD40L mutein Human CD40-L cDNA

> AAQ57984 AAQ9409

1816 1816 840

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CD40 ligand; alteration; immunoreactivity; human cell; accessory molecule ligand; AML; gene therapy; treatment; neoplasia; autoimmune disorder; rheumatoid arthritis; vaccine; chimera; ss.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

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Description	Exemplary CD40 lig Exemplary CD40 lig Human CD40-L type CD40 ligand gene u CD40 ligand coding CD40-L DNA. Homo CD40 ligand gene. Human CD40 ligand cDNA of CD40L, a n
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SUMMARIES	AAV39003 AAV39003 AAV38997 AAV12852 AAV12852 AAQ41506 AAQ4123 AAT05763 AAT93782
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% Query Match Length DB	786 786 786 786 786 840 840 840
8 Query Match	100 .0 98 .8 97 .2 97 .2 97 .2 97 .2 97 .2
Score	786 776.4 763.6 763.6 763.6 763.6 763.6
Result No.	H G W A R D C B Q

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(REGC) UNIV CALIFORNIA. Cantwell M, Kipps TJ, WPI; 1998-348521/30.

97WO-US22740 97US-0982272, 96US-0032145,

08-DEC-1997; 01-DEC-1997; 09-DEC-1996;

Chimeric - Mus sp. . Chimeric - Homo sapiens.

WO9826061-A2 18-JUN-1998.

	18	754 754	96.0 95.0	786	13 -	AAV39000	ΑĞ	Human CD40-L CDNA. Exemplary CD40 liq
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	22	N		906	22	AAF82932	IH .	
	23	642.8	81.8	2209	22	AAF82929	H	HIV-1 gp120-human
	24	$^{\circ}$	81.8	2252	22	AAF82928	H	П
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	56		81.2	1425	14	AAQ41516	Hu	Human CD40-L/Fc fu
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	28	637.2	81.1	929	18	AAT58123	ᄗ	cDNA encoding yeas
	59	637.2	81.1	929	20	AA227537	æ	crimeri
	30	608.4	77.4	882	21	AAZ55540	Fe	CD154
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	32	605.4	77.0	780	21	AAZ55542	Fe	
Ö	33	605.4	77.0	780	21	AAZ55543	Fe	Feline CD154 cDNA
	34	584.4	74.4	783	15	AAQ63960	Œ	Mouse CD40-L type
	35	584.4	74.4	783	19	AAV12853		CD40 ligand coding
	36	584.4	74.4	783	20	AA227524	W	Mouse CD40-L codin
٠	37	582.8	74.1	783	19	AAV38998		
	38	582.8	74.1	818	19	AAV61062	¥.	Murine CD40 ligand
	39	581.2		1878	. 21	AAZ55534	S	CD154
ပ	40	581.2	73.9	1878	21	AAZ5535'	S	Canine CD154 cDNA
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The present sequence represents an exemplary CD40 ligand gene, comprising nuclectides encoding the extracellular domains (Domains III and IV) and transmembrane domain (Domain II) of human CD40 ligand gene (AAV38998) operatively linked to nuclectides encoding the cytoplasmic domain (Domain I) of the murine CD40 ligand gene (AAV38997). The sequence is used to exemplify the method of the invention. The specification comprises a method for altering the immunoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells. Vectors containing the AML genes can be used in gene therapy for treating neoplasia or autoimnume disorders such as rheumatoid arthritis. They can also be used for vaccination to produce immunity against a virus cell, bacteria, protein, fungus or neoplasia.
Vectors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatment of neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
                                                                                                                                                                                                                                                106; 167pp; English
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Sequence 786 BP; 250 A; 166 C; 171 G; 199 T; 0 other;

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Length 786;
                  Indels
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Best Local Similarity 100.0%; Pred. No. 2.4e-219;
Matches 786; Conservative 0; Mismatches 0;
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Conservative

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Matches

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Similarity

Query Match Best Local S

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The present sequence represents an exemplary CD40 ligand gene, comprising nuclectides encoding the extracellular domains (Domains III and IV) of the human CD40 ligand gene (AAV38998) operatively linked to nucleotides encoding the cytoplasmic domain (Domain I) and transmembrane domain (Domain II) of the murine CD40 ligand gene (AAV38997). The sequence is used to exemplify the method of the invention. The specification describes a method. For altering the immunoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells. Vectors containing the AML genes can be used in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              gene therapy for treating neoplasia or autoimmune disorders such as rheumatoid arthritis. They can also be used for vaccination to produce immunity against a virus cell, bacteria, protein, fungus or neoplasia.
                                                                                                                                                                                                                                                                                                                                           neoplasia;
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              GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    vectors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatment neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
                                                                                                                                                                                                                                                                                        Exemplary CD40 ligand gene used in the course of the invention.
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                                                                                                                                                                                                                                                                                                                       CD40 ligand; alteration; immunoreactivity; human cell; accessory molecule ligand; AML; gene therapy; treatment; neogautoimmune disorder; rheumatoid arthritis; vaccine; chimera;
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pred. No. 1.5e-216;
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Chimeric - Homo sapiens.
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                                                            GAAGATTTTGTATTCATGAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC
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                                 CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT
                                                                                                                         Leucine zipper; trimerisation; trimeric CD40-L; fusion protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         hetero-oligomer, homo-oligomer, type II transmembrane protein, soluble CD40-L; tumour necrosis factor family; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                     Human CD40-L type II transmembrane protein coding sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A DNA fragment encoding the extracellular (soluble) region of human CD40-L was ligated to a synthetic oligonucleotide sequence coding for a leader peptide, a 33 amino acid leucine zipper sequence (AAR$3968) and the Flag (RTM) linker sequence. Cells expressing the fusion construct are grown to accumulate oligomeric, soluble CD40-L in the supernatant. The leucine zipper sequence spontaneously trimerises in solution and fusion proteins comprising
/product- human_CD40-L
/note- "nucleotides 148-783 code for the
extracellular region (amino acids 50-261)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAAGATTTTGTATTCATGAAACGATACAGAGATGCÀACACAGGAGAAAGATCCTTATCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the sequence fused to a heterologous mammalian protein also
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Pred, No. 8.3e-213;
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                                                                                                                     TTCGAGAGAATCTTACTCAGAGCTGCAAÄTACCCACAGTTCCGCCAAACĆTTGCGGGCAA
                                                                                                                                  CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                                                                   gctgaaaaaggatactacaccatgagcaacaacttggtaaccctggaaaatgggaaacag
                                      CD40 ligand; alteration; immunoreactivity; human cell; accessory molecule ligand; AML; gene therapy; treatmen autoimmune disorder; rheumatoid arthritis; vaccine; ss
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Length 786;
   Score 763.6; DB 19;
Pred. No. 8.3e-213;
); Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   standard; cDNA to mRNA; 786 BP.
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     97.2%;
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                        Conservative
                Similarity
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781 ctctga 786
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      Query Match
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                                                                     Leucine zipper; fusion protein production; soluble oligomeric protein; heterologous mammalian type II transmembrane protein; activated T cell; heterologous mammalian type II transmembrane protein; antibody production; CD40-L; B-cell proliferation; CD27-L; lymphocyte antigen; ss.
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Pred. No. 8.3e-213;
0; Mismatches 14;
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91US-0805723.
92US-0969703.
93US-0107353.
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CD40 ligand coding sequence:
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Best Local Similarity
Matches 772; Conserv
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05-DEC-1991;
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CTTTTTCCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT
                                                                                                             TTACTGAACTGTGAGGAGTTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA
                                                                                                                                                                               GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG
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                                                                                                                                                                                                                                                                                                            This sequence encodes a human CD40-L polypeptide which binds to CD40. CD40-L is a type II membrane polypeptide which has an extracellular region at its C-terminus, a transmembrane region and an intracellular region at its N-terminus. A soluble form of CD40-L lacks the transmembrane domain. CD40-L activity is mediated by
                                                                                                                                                                                                                                                                                                                                                                            lacks the transmembrane domain. CD40-L activity is mediated by binding with CD40 an dinduces B cell proliferation and induction of antibody secretion, including IgE. Membrane bound CD40-L acts as a CD40 agonist and soluble CD40-L acts as a CD40 antagonist. CD40-L can be used in a binding assay to detect cells expressing CD40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New cytokine CD40-L as CD40 agonist and antagonist - is used for treating allergies, lupus, rheumatoid arthritis, graft-versus-host disease and insulin-dependent diabetes mellitus
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Pred. No. 8.6e-213;
0; Mismatches 14; 1
                                                                                                                                          Spriggs MK;
                                                                                                                                                                                                                                                                                     Claim 1; Fig 2; 80pp; English.
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Local Similarity 98.2%;
les 772; Conservative
                                                             91US-0783707.
                              92WO-US08990
                                                                                                                                             Fanslow WC,
                                                                                                             (IMMV ) IMMUNEX CORP
                                                                                                                                                                           WPI; 1993-152417/18.
                                                                                                                                                                                          P-PSDB; AAR36701
                                                                                                                                             Armitage RJ,
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05-DEC-1991;
                              23-OCT-1992;
29-APR-1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Probe; primer; PCR; amplify; polymerase chain reaction; detection; mutation; CD40 ligand gene; IgM; ss.
                                                                                                                                                                       CGGGAAGCTTCGAGGTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                    TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA

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/product= CD40 ligand
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94US-0184422.
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P-PSDB; AAR57469.
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20-JAN-1994;
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TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
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                                                                                                                                                                                                                             ;
                                         840;
                                          Length
                                                          Indels
                 214 T; 0 other;
                                        Score 763.6; DB 15;
Pred. No. 8.6e-213;
0; Mismatches 14;
                C; 175 G;
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                                         97.2%;
98.2%;
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               840 BP; 266 A; 185
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                                                Local Similarity 98.2 les 772; Conservative
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hyperigm syndrome
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                 Sequence
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                                         Query Match
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14;

0; Mismatches

Length

Score 763.6; DB 16; Pred. No. 8.6e-213;

97.2%;

other;

BP; 263 A; 182 C; 181 G; 214 T; 0

240 261

201

141

321 360

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high-density
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            the
producing
                                                       B-cell;
frugiperda;
                                                    High density membrane.bound CD40 ligand; B-lymphocyte, B-cell, differentiation; proliferation; baculovirus; Spodoptera frugis Sf9; insect cell culture; tumour necrosis factor receptor; ss

    for stimulating
vivo, partic. for

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CD40 ligand
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us-08-982-272-6.rng

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GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780
                                                                                                       CGGGAAGCIICGAGICAAGCICCAITIAIAGCCAGCCICIGCCIAAAAGICCCCCGGIAGA 600
                                                                                                                                           099
                                                                                                                                                             681
                                                                                                                                                                                                741
                                                                                                                                                                                                                          GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG 480
                                                                              immune deficiency;
                                                                                                                                                                                                                                                                                                                                                                                    monoclonal antibody;
                                                                                                                TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                               CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                                                                                                                                                                        CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Treating or preventing neoplastic disease associated with CD40-expressing cells - particularly B-cell lymphoma, by administration of CD40-binding protein, preferably antibody soluble CD40-ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WJ;
                                                                                                                                                                                                                                                                                                                                                                                    Cytokine; activated CD4+ T cell; CD40; CD40L; mo neoplastic cell proliferation; B-cell lymphoma; melanoma; carcinoma; ss.
                                                                                                                                                                                                                                                                                                                                                                    cDNA of CD40L, a novel cytokine ligand for CD40.
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers 46..831
                                                                                                                                                                                                                                                                                                                   AAT93782 standard; cDNA; 840
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(USSH ) US DEPT HEALTH
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The present sequence represents the CDNA sequence of a novel cytokine ligand for CD40 called CD40L is a type II membrane polypeptide, which is expressed by activated CD4+ T cells, and causes B cell proliferation and induction of antibody secretion. The protein can be used to produce monoclonal antibodies, which in turn bind to CD40-expressing cells. This inhibits binding of soluble CD40 to its proliferation of neoplastic cells, and is particularly useful in treating proliferation of neoplastic cells, and is particularly useful in treating proliferation of e.g. where induced after transplants or in other cases of immune deficiency such as ADDS), and also melanoma or carcinoma. Since the monoclonal antibodies inhibit neoplastic cells directly, they may not
                                                                                                                                                                                                                                                                                                                    octonal antibodies inhibit neoplastic cells directly, they may not be coupled to a toxin or radioisotope, avoiding toxic effects on
                                                                                                                                                                                                                                                                                                                                                                                                                              · Sequence 840 BP; 266 A; 185 C; 175 G; 214 T; 0 other;
                Claim 3; Columns 19-22; 21pp; English.
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721 GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780 540 645 465 480 525 225 240 285 300 345 405 420 ATGAAGATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120 180 Gaps ttogagagaatettaetteagagetgeaaataeceacagtteegeeaaeettgegggeaa CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT gctgaaaaaggatactacaccatgagcaacactggtaaccctggaaatgggaaacag GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTAACAGTGG GCTGAAAAAGGATACTACACCATGAGCAACATTGGTAACCCTGGAAAATGGGAAACAG 1 ATGATAGAAACATACAGCCAAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC ö Length 840; Indels Score 763.6; DB 18; Pred. No. 8.6e-213;); Mismatches 14; 97.2%; 98.2%; Conservative Similarity Query Match Best Local Simi Matches 772; 646 466 526 541 586 601 421 181 166 181 226 241 361 61 121 qq g δ Q δ ŏ Dp, q p, ŏ ΟŽ g ò qq QY ò g οy δ

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence encodes human CD40 ligand which is used in the method of the invention. The method has been developed for proliferating be cells to increase their number at least 100-fold. The method comprises: (a) providing high density, membrane bound CD40 ligand; and (b) culturing one or more B cells in the presence of this ligand. The culture results in a proliferation in the number of B cells of at least 100 fold. Also described is a method as above where the B cells are induced to differentiate into antibody-producing cells in the presence of one or more cytokines. The method can be used for stimulating B-cell proliferation and activation is suppressed. Eight rounds of division over six days can be achieved, corresponding to a 256-fold
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             numbers, which is a vast increase compared to previous
          766 gigacigatecaagecaagigagecaiggeactggeticaegiectitggetiaeteaaa 825
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   sed proliferation of B cells in culture - by incubating them presence of membrane-bound CD40 ligand
                                                                                                                                                                                                         Human; CD40 ligand; TNF receptor family; activated T cell;
type 2 membrane glycoprotein; cell proliferation; differentiation;
Sequence 840 BP; 263 A; 182 C; 181 G; 214 T; 0 other;
                                                                                                                                                                                   Human CD40 ligand encoding DNA sequence.
                                                                                                                                                                                                                                                                                                 /*tag= a
/product= "CD40 ligand"
                                                                                                                                                                                                                                                                                                                                                                                                                                      (BOEH ) BOEHRINGER INGELHEIM PHARM INC.
                                                                                                                                                                                                                                                                           Location/Qualifiers
22..807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 1; Fig 1; 37pp; English.
                                                                                                               AAV61063 standard; DNA; 840 BP.
                                                                                                                                                                                                                                                                                                                                                                                                     95US-0431055
94US-0234580
                                                                                                                                                                                                                                                                                                                                                                               95US-0431055
                                                                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kehry M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          culturing methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cell
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                                                       ctctga 831
                                 CTCTGA 786
                                                                                                                                                                                                                                                      Homo sapiens
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28-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                               8-APR-1995;
                                                                                                                                                             08-DEC-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               increase in
                                                                                                                                                                                                                                                                                                                                  US5817516-A
                                                                                                                                                                                                                               cell; ds
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Castle B,
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                                                                                                                                      AAV61063;
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                                                                                                      AAV61063
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CD40-L; CD40 receptor ligand; anti-human CD40-L monoclonal antibody; binding inhibitor; trimeric CD40-L; anti-Immunoglobulin M; peripheral blood B cell; proliferation inhibitor; ss.
                                                                 GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACAAGAGAAAGATCCTTATCC
                                                                                                                                                                                                                                                                                                                               ATGAAGATTTTTATGTATTTACTTACTGTTTTTTTTTTATCACCCAGATGATTGGGTCAGCA
                                                                                                                  CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGAAGGAATCTTCAT
                                                                                                                                                                                                                  gaagattttgtattcatgaaaacgatacagagatgcaacacaggagaaagatccttatcc
                                                                                                                                                                                                                                                                                                                AACAAAGAGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                                                                                                                                                                                                                                               CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
                                                                                                                                                                                                                                                                                                                                                                                               CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         562 cgggaagcttcgagtcaagctccatttatagccagcctctgcctaaagtcccccggtaga
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ttcgagagaatcttactcagagctgcaaatacccacagttccgccaaaccttgcgggcaa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
                                                                                                                                                                                                                                                  TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA
                                                                                                                                                                                                                                                                                                                                                                                                                                             GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              standard; cDNA; 840
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human CD40-L coding sequence
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Gaps

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14; Indels

Length 840;

DB 19;

Score 763.6; DB 19. Pred. No. 8.6e-213; 0; Mismatches

97.28; 98.28;

Best Local Similarity 50.2 Matches 772; Conservative

Query Match

705

-08-982-272-6.rng

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The present sequence encodes a qp39 protein. It is a CD40 ligand. CD40 ligands are used for reducing osteoblast cell death or apoptosis, and for treating or preventing bone loss in animals, preferably humans, at risk of, or undergoing, bone loss. The bone loss is associated with osteoporosis, osteonecrosis, inflammatory arthritis, post-menopausal osestrogen loss, esterogen loss due to ovariectomy, total historectomy, lupus nephritis, Takayasu's arteritis; Wegener's granulomatosis, anti-glomerular basement membrane nephritis, myositis, scleroderma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           apoptosis associated with osteoporosis, osteonecrosis and inflammatory arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 idiopathic autoimmune thrombocytopenia, asthma, a chronic obstructive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phrotic/nephritic syndrome, or cancer. They may also prevent bone loss in a subject undergoing, or schedu
                                                                                                                                                                                                                                                                                                                                                                                     9939; CD40 ligand; osteoblast cell death; apoptosis; bone loss; osteoporosis; osteonecrosis; inflammatory arthritis; esterogen los ovariectomy; historectomy; lupus nephritis; Takayasu's arteritis; Megener's granulomatosis; nephritis; myositis; scleroderma;
                                                                                                                  GIGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
                                                                                                                                 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
               caatccattcacttgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaat
                                                        CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                                                                                                                                                                                                                                                                                                                                             Nucleotide sequence of human gp39 protein, a CD40 ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                   thrombocytopenia; asthma; lung disease; cancer; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 113; 118pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
22.807
/*tag= a
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to treat or prevent bone loss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "gp39"
                                                                                                                                                                                                                                                                                  standard; DNA; 879 BP.
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                                                                                                                                                                                   781 CTCTGA 786
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                                                                                                                                                                                                                826 ctctga 831
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                                                                                                                                                                                                                                                                                   This sequence encodes the human CD40 receptor ligand (CD40-L). The invention relates to anti-human CD40-L monoclonal antibodies M90 secreted by hybridoma hCD40L-M90 (ArCC HB-12055) and M91 secreted by hybridoma hCD40L-M91 (ArCC HB-12055). M90 and M91 inhibit CD40-L binding to CD40 and the ability of trimeric CD40-L and anti-Immunoglobulin M to induce proliferation of peripheral blood B cells.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATGAAGATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 763.6; DB 20;
Pred. No. 8.6e-213;
0; Mismatches 14;
                                                                                                                                                                                                                                       Anti-human CD40-Ligand monoclonal antibodies
                                                                                                                                                                Armitage RJ;
                                                                                                                                                                                                                                                                 Disclosure; Fig 2; 59pp; English;
                                                                                                                                                                                                                                                                                                                                                                                                                                       97.2%;
98.2%;
                                          94US-0249189
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92US-0969703.
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                                                                                                                                       (IMMV) IMMUNEX CORP.
                                                                                                                                                                    Fanslow
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Matches 772; Conserv
                                                                                                                                                                                                            P-PSDB; AAY39938
                                             24-MAY-1994;
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                                                                            25-OCT-1991;
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                   05-0CT-1999
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                                                                                                                            TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA
                                                                                                                                                                                    1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC
                                                                           atgatcgaaacatacaaccaaacttctccccgatctgcggccactggactgcccatcagc
                                                                                                                    CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGAAGGAATCTTCAT
                                                                                                                                                 GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC
                                                                                                                                                        AACAAAGAGAGAGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                                                                               caaattgcggcacatgtcataagtgaggccagcagtaaaacaacatctgtgttacagtgg
                                                                                                                                                                                                                                                             GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG
                                                                                                                                                                                                                                                                    CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                                                                                                                                                    CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
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             C; 190 G; 222 T; 0 other;
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                                 Score 763.6; DB 22
Pred. No. 8.8e-213;
                                               0; Mismatches
bone marrow transplant.
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98.28;
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CD40 engagement on T cells can be used to induce T cell receptor rearrangement and enhance T cell affinity for a particular antigen. CD40 engagement can be bought about by contacting CD40 with a CD40-binding agent, e.g. human CD40 ligand. The CD40-binding agents can also be used in methods for promoting T cell maturation, inhibiting T cell receptor rearrangement, inhibiting environmental stress-induced cell death, altering the specificity of a T cell towards an antigen or enhancing environmental stress-induced cell death (all claimed). T cell affinity maturation towards a specific antigen can be inhibited, especially for a self-antigen in an autoimmune disease, which includes rheumatoid arthritis, uveitis, insulin dependent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        mellitus, haemolytic anaemias, rheumatic fever, Crohn's disease.
Barre syndrome, psoriasis, thyroiditis, Grave's disease,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Guillain-Barre syndrome, psoriasis, thyroiditis, Grave's disease, myasthenia gravis, glomerulonephritis, autoimmune hepatitis or systemic lupus erythematosus. Inducing environmental stress-induced T cell death is carried out in a cancerous T cell or a self-reactive T cell where the
                                                    CD40 ligand; CD40; T cell; T cell receptor; rearrangement; maturation; cell death inhibition; stress-induced; immunosuppressive; anti-thyrold; anti-inflammatory; anti-diabetic; anti-rheumatic; anti-anaemic; opthalmological; anti-psoriatic; nephrotophic; hepatotropic; virucide; dermatological; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            219
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        T cell receptor gene rearrangement for treating autoimm comprises contacting T cells with a CD40-binding agent
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Pred. No. 1.2e-212;
0; Mismatches 14; 1
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         GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC 240
                                   TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
                                            Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder; ss.
                                                                                                                                                                                                                                                                                                                                       TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ONA encoding novel human diagnostic protein #22375.
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23-AUG-2000;
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The invention relates to isolated polynucleotide (I) and collaboration rescues. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome cand gene mapphig, and in recombinant production of (II) The collaboration of collaboration of collaboration of collaboration of collaboration of collaboration and in recombinant production of (II) The collaboration of colla
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                                                New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
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Pred. No. 1.2e-212;
0; Mismatches 14;
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                                                                                                                                                                        Claim 1; SEQ ID No 22375; 103pp; English
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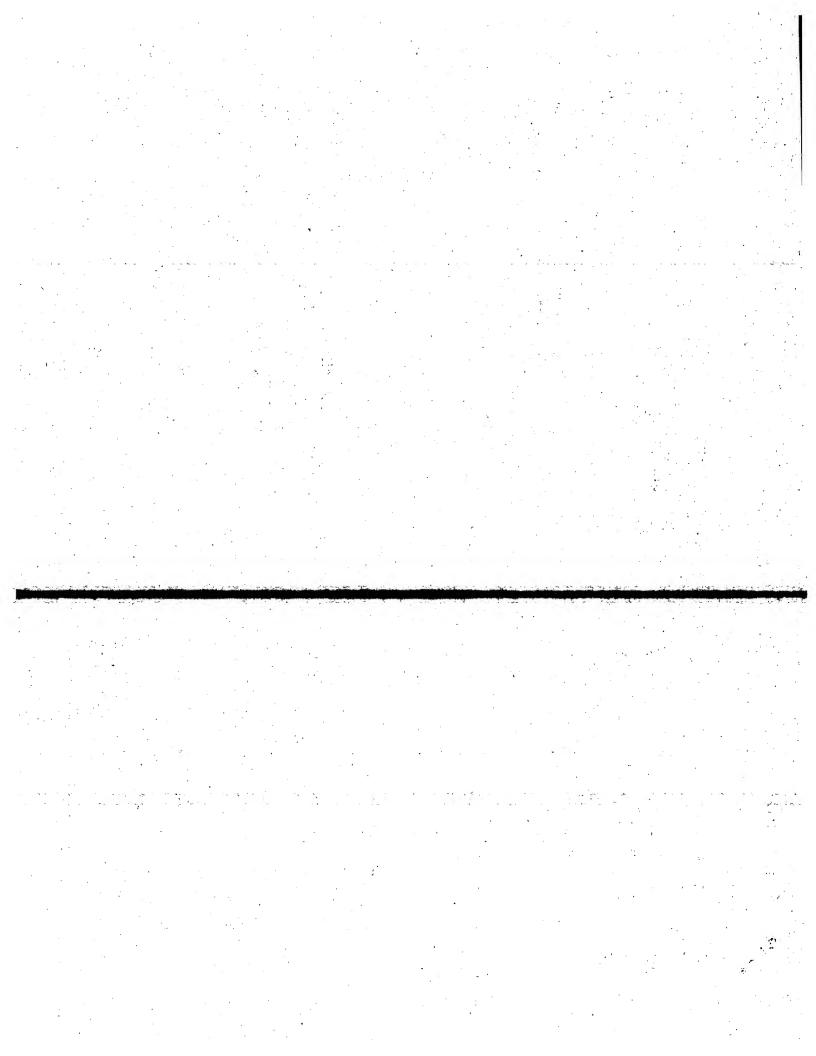
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                                                                                                               caatccattcacttgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaat 759
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                 GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
                                                                                            CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                                                                                     which is
                                                                                                                                                                                                                                                                                                             gp39; T-cell antigen; CD40 ligand; B-cell proliferation;
B-cell differentiation; ss.
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ligand for the CD40 receptor, causing proliferation and
differentiation of B cells and some cancer cells
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P-PSDB; AAR59548.
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Length 840;
                     Indels
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2.5e-212;
                     Mismatches
 Score 762;
            Pred. No.
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Job time: 11260 sec
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(without alignments)
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1 ATGATAGAAACATACAGCCA......TTGGCTTACTCAAACTCTGA 786
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd
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Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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ptodata/2/pna/US6021_COMB. ptodata/2/pna/US6022_COMB. ptodata/2/pna/US6023_COMB.

'2/pna/US6017

/pna/US6027

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

•	Description	Sequence 6, Appli	,	3, 7	0	Sequence 7, Appli	1,	'n	10,	11,	11,	7, A	ì	'n	m	Sequence 1, Appli	11,	11,	4,7	1, A	11,	Sequence 11, Appl	11,		٦	11,	1, A	Sequence 1, Appli	٦,	11,	Sequence 11, Appl	Sequence 1, Appli
	ID	US-08-982-272-6	US-08-982-272-7	US-08-107-353-3	US-08-982-272-1	PCT-US94-00786-7	PCT-US94-14767-1	PCT-US97-11956-1	-802-7	US-07-969-703A-11	-696-	US-08-009-258-7	US-08-172-664-1	215-	234-5	96-2	US-08-477-733A-11	US-08-484-624-11	US-08-589-771-7	US-08-677-762-1	US-08-770-981-11	088-913	322-02	09-322-	09-365-940-		US-09-387-935-1	-60-	09-388-07	09-392-61	-09-399-10	US-09-430-448-1
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MODULATION AND TREATMENT
MALIGNANCIES
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VENTION: NOVEL EXPRESSION VECTORS
  Mismatches
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GENERAL INFORMATION:
APPLICANT: Kipps, Thomas J.
APPLICANT: Sharma, Sanjai
     Matches '786; Conservativé
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TITLE OF INVENTION: OF TITLE OF INVENTION: OF TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION: NUMBER OF SEQUENCES:
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Sequence 31, Appli
Sequence 31, Appli
Sequence 3, Appli
Sequence 22375, A
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Sequence 6,
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Pred. No. 1.8e-201;
           US-09-545-993-7
US-09-545-9938-7
US-09-645-9938-7
US-09-645-9268-1
US-09-909-595-3
US-09-909-595-3
US-00-329-393-3
PCT-US01-08631-22375
PCT-US01-0865-373
PCT-US99-3030-1
US-09-442-384A-457
US-09-470-494-1
US-09-470-494-1
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                                                                                                                                                                                                                                                                                                                                                                                 CONTAINING ACCESSORY
MOLECULE LIGAND GENES AND
THEIR USE FOR IMMUNO-
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                                                                                                                                                                                                                               ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER REDABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: 1.44 Mb storage
COMPUTER: 1BM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSeq Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/982,272
FILING DATE:
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ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                           RESULT 1
US-08-982-272-6
; Sequence 6, Application US/08982272
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Guise, Jeffrey W. REGISTRATION NUMBER: 34,613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
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ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                            Kipps, Thomas J.
                                                                                                                                                                                                                                                                                                                                            Sharma, Sanjai
Cantwell, Mark
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TYPE: nucleic acid
STRANDEDNESS: single
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 6
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Los Angeles
California
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US-08-982-272-6
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APPLICANT:
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pred. No. 7.1e-199;
0; Mismatches 6;
                                                                                                                                                                                                                                US/08/982,272
                                                                                                                                                                              OPERATING SYSTEM: IBM P.C. DOS SOFTWARE: FastSeq Version 2.0
                                                                                                                                                                                                                                                  FILING MALE TOWN: CLASSIFICATION: PRIOR APPLICATION DATA:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                              MEDIUM TYPE: 3.5" Diskette, MEDIUM TYPE: 1.44 Mb storage
                                                                                                                                                                                                                                                                                                                                                   NAME: Guise, Jeffrey W. REGISTRATION NUMBER: 34,613
                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                         (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (213) 955-0440
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 786 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            98.8%;
99.2%;
                                                                                                                                                                 IBM Compatible
                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 60/13
FILING DATE: 12/9/96
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 99.2'
Matches 780; Conservative
                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE:
                                                                                                                COMPUTER READABLE FORM:
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US-08-982-272-7
                                                                                RY: U.S.A.
90071-2066
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                                                                                                                                                                 COMPUTER:
                                                                               COUNTRY:
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                                                                                       CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                                                                                                                                                          TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
TITLE OF INVENTION: Proteins
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/107,353
FILING DATE: 19930813
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1: Immunex Corporation
51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 3, Application US/08107353 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 786 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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LOCATION: 1...
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US-08-107-353-3
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                                                                                                                    ATGAAGATTTTTATGTATTTACTTACTGTTTTTTTTTATCACCCAGATGATTGGGTCAGCA 120
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                                                                       1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC 60
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                           Length 786;
                                                   Indels
                           Score 763.6; DB 5;
Pred, No. 2.1e-195;
0; Mismatches 14;
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GENERAL INFORMATION:
APPLICANT: Kipps, Thomas J.
APPLICANT: Sharma, Sanjai
"SEPLICANT: Cantwell, Mark
TITLE OF INVENTION: NOVEL EXPRESSION VECTORS
TITLE OF INVENTION: CONTAINING ACCESSORY
                                97.2%;
98.2%;
                                                        Conservative
                              Query Match
Best Local Similarity
Matches 772; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CTCTGA 786
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US-08-107-353-3
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98.2%; Pred. No. 2.1e-195;
tive 0; Mismatches 14;
        THEIR USE FOR IMMUNO-
MODULATION AND TREATMENT
                                                                                                                                                                                                                                                                                                         US/08/982,272
                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: 1.44 Mb storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS
                                                                                                                                                                                                                                                                 SUBTWARE: FastSeq Version 2.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/9R7
                                        MALIGNANCIES
                                                                                                                                                                                                                                                                                                                                                                          60/132145
                                                                                          : Lyon & Lyon
633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Guise, Jeffrey W. REGISTRATION NUMBER: 34,613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
60/132
FILING DATE: 12/9/96
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              786 base pairs
nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                         CORRESPONDENCE ADDRESS:
                          INVENTION: P
                                                                                                                      Suite 4700
                                                                                                                                       Los Angeles
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 772; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (213)
                                                                                                                                                                   COUNTRY: U.S.A.
ZIP: 90071-2066
               INVENTION
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US-08-982-272-1
                                                                                           ADDRESSEE
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                                                                     CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCCAAGTCACCTTCTGTTCCAAT
                                                                                   TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                  TITLE OF INVENTION: DETECTION AND TREATMENT OF MUTATIONS TITLE OF INVENTION: IN A CD40 LIGAND GENE NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: MS Word for Apple 5.1, Version a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCT/US94/00786
                                                                                                                                                                                                                                                                                                                                                                                            Sequence 7, Application PC/TUS9400786 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                          FANSLOW, WILLIAM
RENSHAW, BLAIR
SPRIGGS, MELANIE
WIDMER, MICHAEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Floppy disk
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PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy
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STATE: WASHINGTON
COUNTRY: USA
ZIP: 98101
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Mismatches
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Pred. No. 2.1
                                                                                                                                                                                                                                               97.2%;
nilarity 98.2%;
Conservative 0
                                                                                                  ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
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772; Conser
                                                                 ANTI-SENSE: NO ORIGINAL SOURCE:
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PCT-US94-00786-7
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Best Local Si
Matches 772
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661 CAATCCATTCACTTGGGAGGAGTATTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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                                    166 CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT
121 CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT
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GENERAL INFORMATION:
APPLICANT: Immunex Corporation
TITLE OF INVENTION: METHOD OF ACTIVATING DENDRITIC CELLS
NUMBER OF SEQUENCES: 4
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SOFTWARE: Microsoft Word for Macini
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US97/11956
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ADDRESSEE: IMMUNEX C
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CITY: SI
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                                                                                                                                                                                        METHOD OF PREVENTING OR TREATING DISEASE CHARACTERIZED BY NEOPLASTIC CELLS EXPRESSING CD40
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                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: Apple Macintosh System 7.1 SOFTWARE: Microsoft Word for Macintosh, Version #5.1a CURRENT APPLICATION DATA:
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pred. No. 2.1e-195;
0; Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRICE APPLICATION DATA:
APPLICATION NUMBER: USSN 08/172,664
FILING DATE: December 23, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATION NUMBER: PCT/US94/14767
DATE: December 21, 1994
                                                                                                          Sequence 1, Application PC/TUS9414767
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                        3E: IMMUNEX CORPORATION -51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY AND ATTORNEY PATTICIA A. REGISTRATION NUMBER: 34,693 REFERENCE/FOOCKET UNBER: 28: TELECOMMUNICATION INFORMATION: TELECHONE: (206)587.9430
                                                                                                                                                                     APPLICANT: LONGO, DAN L.
APPLICANT: MURPHY, WILLIAM
TITLE OF INVENTION: METHOD OF
TITLE OF INVENTION: DISEASE C
TITLE OF INVENTION: EXPRESSIN
NUMBER OF SEQUENCES: 2
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                                                                                                                                           APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                              FANSLOW, WILLIAM
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MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
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SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
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IMMEDIATE SOURCE:
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46..831
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Best Local Simi
Matches 772;
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CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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721 GIGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
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COMPUTER: IBM PC COMPATIBLE
OPERATORS SYSTEM: PC-DOS/MS-DOS
SOFTWARRE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 763.6; DB 3;
Pred. No. 2.1e-195;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/805,72
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SPRIGGS, MELANIE
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                     Sequence 10, Application US/07805723 GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD APPLICANT: FANGION, WILLIAM ANDLICANT: FANGION, WILLIAM ANDLICANT: ANDLICANT.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: IMMUNEX CORPORATION 51 UNIVERSITY STREET
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NAME: OSTER, JEFFREY B.
REGISTRATION NUMBER: 32895
REFRENCE/CDOCKET NUMBER: 280
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 2065870606
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
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98.2%;
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Best Local Similarity 98.2*
Matches 772; Conservative
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COMPUTER READABLE FORM:
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US-07-805-723-10
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US-07-805-723-10
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Pred. No. 2.1e-195;
0; Mismatches 14;
                USSN 08/677,762
                                                                   USSN 08/763,995
                                                                                                               NAME: Perkins, Particia A.
REGISTATION NUMBER: 34,693
REPERENCE/DOCKET NUMBER: 2845:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEPHONE: (206)233-0644
INFORMATION FOR SED ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              97.2%;
98.2%;
                           FILING DATE: 10-JUL-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN
FILING DATE: 12-DEC-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                               LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
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Best Local Similarity 98.2
Matches 772; Conservative
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PRIOR APPLICATION DATA:
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LOCATION:
PCT-US97-11956-1
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                                                                 CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180.
                                                                            GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC 240
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                                      TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
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                                                                                                                                                                                                                                                                                                        NOVEL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 11, Application US/07969703A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: ARMITAGE, RICHARD APPLICANT: FANSLOW, WILLIAM APPLICANT: SPRIGGS, MELANIE ATTLE OF INVENTION: NOVEL CYTON NUMBER OF SEQUENCES: 17 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5: IMMUNEX COR
51 UNIVERSITY
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|CTCTGA 831
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CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT

ATGAAGATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA

1 ATGATAGAAACATACAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC. 60

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Length 840; Indels

97.2%; Score 763.6; DB 3; 98.2%; Pred. No. 2.1e-195; ive 0; Mismatches 14;

Query Match 97.2 Best Local Similarity 98.2 Matches 772; Conservative

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SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/969.7022
COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: PERKINS, PALTICIA A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                                                                                                                                                                                                                                       LENGTH: 840 base pairs
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                        CLASSIFICATION:
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                                                                                          FILING DATE:
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US-07-969-703A-11
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Query Match
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                                                                               CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
                                                                                                                                     CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 600
                                     586 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 645
                                                               TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
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                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                            RESULT 10
US-07-969-703B-11
; Sequence 11, Application US/07969703B
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEE: IMMUNEX CORPORATION: 51 UNIVERSITY STREET
SEATTLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 26
TELECOMUNICATION INFORMATION
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
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NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,6
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TYPE: NUCLEIC ACID
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US
FILING DATE: 19921023
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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; LOCATION: 46..831
US-07-969-703B-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: HOMO
IMMEDIATE SOURCE:
CLONE: CD40-L
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
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                                                                                                                                                                                                                                       781 CTCTGA 786
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                                                                                            46 ATGATCGAAACATACAACCATCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 105
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                                                                            1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC
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  Length 840;
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Score 763.6; DB.3;
Pred. No. 2.1e-195;
); Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: WIDMER, MICHAEL
TITLE OF INVENTION: DEPECTION AND TREATMEN
TITLE OF INVENTION: IN A CD40 LIGAND GENE
NUMBER OF SEQUENCES: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 7, Application US/08009258 GENERAL INFORMATION:
  97.28;
98.28;
                                       Conservative
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                    Similarity
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826 CTCTGA 831
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US-08-009-258-7
                      Best Local Sir
Matches 772;
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CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT -540
                                                                               541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 600
                                                                                                                                                                                             TICGAGAGAAICITACICAGAGCIGCAAATACCCACAGIICCGCCAAACCIIGCGGGCAA 660
GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG 480
                                                                                                                                                                                                                                                         661 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                                                                                                                                                                                                                                                                                                         GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
                 APPLICANT: LONGO, DAN L.
APPLICANT: MURPHY, WILLIAM
TITLE OF INVENTION: METHOD OF PREVENTING OR TREATING
TITLE OF INVENTION: DISEASE CHARACTERIZED BY NEOPLASTIC CELLS
TITLE OF INVENTION: EXPRESSING CD40
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM: Apple Macintosh System 7.1 SPETAMENG SYSTEM: Apple Macintosh, Vers: CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/172,664
FILING DATE: December 23, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: IMMUNEX.CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/08172664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WILLIAM
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nucleic acid
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US-08-172-664-1
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTTTTTGCTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
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pred. No. 2.1e-195;
0; Mismatches 14;
                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/009.258
                                                                                                                                                                                                      19930122
                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                         : IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                NAME: PERKINS, PATRICIA A. REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 281 TELECOMMUNICATION INFORMATION:
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98.2%;
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IMMEDIATE SOURCE:
CLONE: CD40-L
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Best Local Similarity 98.2
Matches 772; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                    840 base pairs
                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                           COMPUTER READABLE FORM:
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                                                                           STATE: WASHINGTON
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                                                          SEATTLE
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                                                                                                                             Gaps
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                                                                                                                                                 1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC
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                                                                                                       Length 840;
                                                                                                                             Indels
                                                                                                       Score 763.6; DB 5;
Pred. No. 2.1e-195;
0; Mismatches 14;
                                                                                                      97.2%;
98.2%;
                                                                                                      Query Match 97.2
Best Local Similarity 98.2
Matches 772; Conservative
                                                   NAME/KEY: CDS
LOCATION: 46..831
          ORGANISM: HOMO
IMMEDIATE SOURCE:
CLONE: CD40-L
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTCTGA 786
                                                              ;
US-08-172-664-1
                                          FEATURE:
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                                                      Hodgkin's
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                                APPLICANT: Gruss, Hans-J_rgen
TITLE OF INVENTION: Method of Diagnosing or Treating
TITLE OF INVENTION: Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 763.6; DB 6;
Pred. No. 2.1e-195;
0; Mismatches 14;
                                                                                                                                                                                                                                                                         OPERATING SYSTEM: Apple Macintosh System SOFTWARE: Microsoft Word for Macintosh, V CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/215,862
                                                                                                CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
Sequence 1, Application US/08215862 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,69 REFERENCE/DOCKET NUMBER: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 97.2%;
98.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 840 base pairs
TYPE: nucleic acid
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Best Local Similarity 98.2
Matches 772; Conservative
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                                                                                                                      ADDALS
STREET: 51 C.
CITY: SEATTLE
THE: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION: 424
                                                                                    NUMBER OF SEQUENCES:
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MOLECULE TYPE: CDN
HYPOTHETICAL: NO.
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ORGANISM: HOI
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US-08-215-862-1
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RESULT 13 US-08-215-862-1

E: DNA CDS 22..807

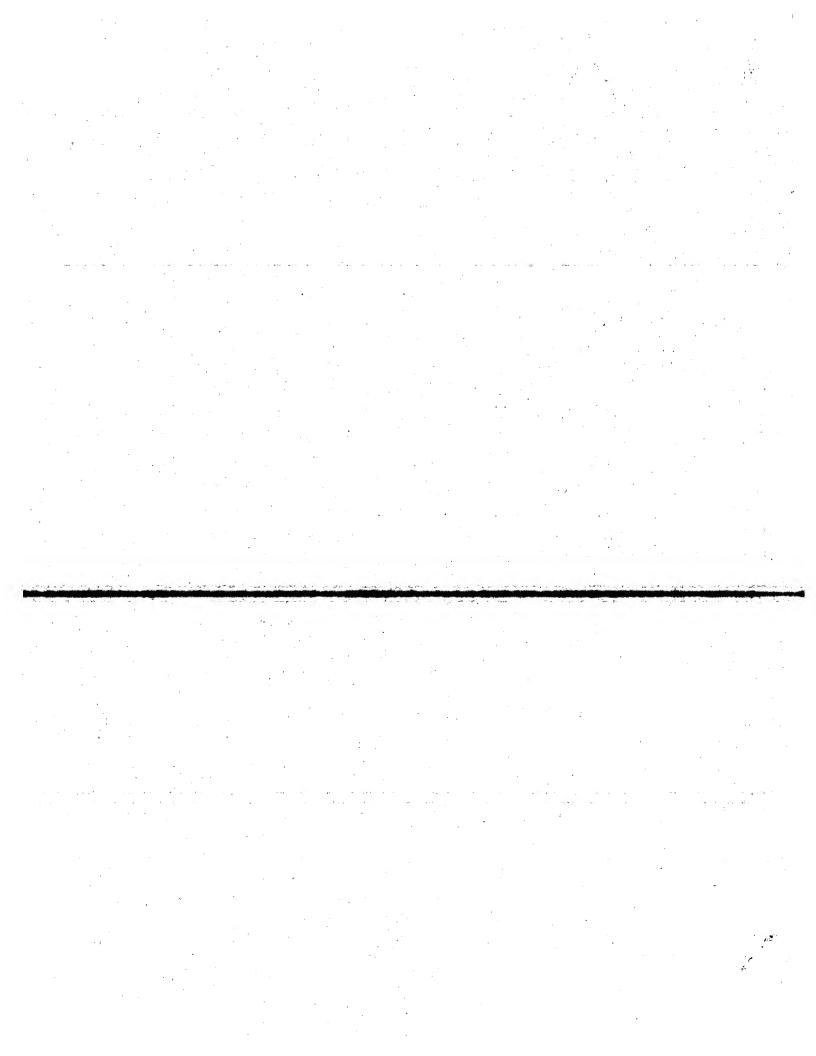
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MOLECULE TYPE:
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                   NAME/KEY:
                             , LOCATION:
US-08-234-580-3
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                                                                                     CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
   USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 3, Application US/08234580
GENERAL INFORMATION:
APPLICANT: KEHRY, MERILYN R
APPLICANT: CASTLE, BRIAN E
TITLE OF INVENTION: METHODS FOR PROLIFERATING AND
TITLE OF INVENTION: DIFFERENTIATING B CELLS, AND U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: MILLMAN, ROBERT A
REGISTRATION NUMBER: 36,217
REFERENCE/DOCKET NUMBER: 1011.1030000/RAM
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            3: STERN, KESSLER, GOLDSTEIN & FOX 100 NEW YORK AVE. N.W. SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC COMPATIBLE OPERATING SYSTEM: PC-DOS/MS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (202)371-2600
TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
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ADDRESSEE: STERN, KE
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US-08-234-580-3
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                                                                                                    CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
                      Gaps
                                                        CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
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                       Indels
Score 763.6; DB 6;
Pred. No. 2.1e-195;
); Mismatches 14;
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US-08.396-230-1
Sequence 1, Application US/08396230
GENERAL INFORMATION:
   97.28;
                            Conservative
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                 Similarity
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       Query Match
Best Local Sim
Matches 772;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
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                            TITLE OF INVENTION: METHOD FOR STIMULATING A CELL-MEDIATED TITLE OF INVENTION: IMMUNE RESPONSE WITH A CD40 BINDING TITLE OF INVENTION: PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM: Apple Power Macintosh System 7.5.5 SOFTWARE: Microsoft Word for Macintosh, Version 6.0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 763.6; DB 7;
Pred. No. 2.1e-195;
0; Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 7;
                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/396,230 FILING DATE: March 1, 1995
                  CHARLES R
                                                                                                                       E: IMMUNEX CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION
TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       97.2%;
98.2%;
                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (206)233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 98.2
Matches 772; Conservative
                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                  MALISZEWSKI
                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                                                                                                                              424
                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                           SEATTLE
WASHINGTON
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                                                                                                                       ADDRESSEE:
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APPLICANT:
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FEATURE:
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721 GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780
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                                                           CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
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Search completed: May 30, 2002, 05:23:08 Job time: 18667 sec



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1 ATGATAGAAACATACAGCCA......TTGGCTTACTCAAACTCTGA 786
                                                                                                         May 30, 2002, 05:39:58 ; Search time 627.19 Seconds
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Compugen Ltd.
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Pending_Patents_NA_New:*

/cgn2_6/ptodata/1/pna/PCT_NEW_COMB.seq:*

/cgn2_6/ptodata/1/pna/USO6_NEW_COMB.seq:

/cgn2_6/ptodata/1/pna/USO8_NEW_COMB.seq:

/cgn2_6/ptodata/1/pna/USO8_NEW_COMB.seq:

/cgn2_6/ptodata/1/pna/USO8_NEW_COMB.seq: /cgn2_6/ptodata/1/pna/US10_NEW_COMB.seq: /cgn2_6/ptodata/1/pna/US60_NEW_COMB.seq:

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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Gaps

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Length 1816;

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1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC

Query Match 97.2%; Score 763.6; DB 5; Length Best Local Similarity 98.2%; Pred. No. 9e-195; Matches 772; Conservative 0; Mismatches 14; Indels

CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180

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241 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300

GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC 240

339

CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 420

301 AACAAAGAGGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 360 aacaaagagagagaagaaagaaagaaacagctttgaaatgcaaaaaggtgatcagaatcct 399

340 361

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	G	226, App	457, App	821, App	131, App	9, Appli	202, App		664, App	7000, Ap	14649, A	35707, A	62305, A	297617,	206146,	206147,	120, App		26090, A	1996, Ap	239, App	1323, Ap	628, App	83794, A	83795, A.	824, App	11996, A
٠	Description	Sequence	Sequence					Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sednence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence	Sequence
SUMMARIES	ID	US-09-053-375B-226	US-09-442-384B-457	US-09-053-375B-821	US-09-911-904-131	US-09-875-453A-9	PCT-US02-10421-202	US-10-112-699-202	US-09-919-002-664	US-09-540-210B-7000	US-09-540-210B-14649	US-10-027-632-35707	US-10-027-632-62305	US-10-027-632-297617	US-10-027-632-206146	US-10-027-632-206147	US-10-123-155-120	US-10-027-632-179264	US-09-539-331D-26090	US-10-105-299-1996	US-10-106-698-239	US-09-789-189-1323	US-09-442-384B-628	US-10-027-632-83794	US-10-027-632-83795	US-10-105-299-824	US-09-919-002-11996
	Query Match Length DB	316 5	316 5	1250 5	508 5	395 5	569 1	9 699	2127 5	234 5	330 5	620 6	520 6	520 6	9 6 9	9 699	775 6	223197 6	438 5	1751 6	1751 6	418 5	10351 5	621 6	621 6	803 6	646 5
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	5 6	34.6	4.4	576	י פ	US-09-721-544-732		732, App	
	30	34.6	4.4	909	٩	US-10-027-632-84272		84272, A	
	31	34.6	4.4	909	9	US-10-027-632-296495		296495,	
	32	34.6	4.4	624	9	US-10-027-632-208277	Sequence	208277,	
	33	34.6	4.4	663	9	-23002	Sequence	230025,	
	34	34.6	4 . 4	1225	9	US-10-027-632-208276		208276,	
υ	35	34.4	4.4	502	'n	US-09-789-189-648		648, App	
	36	34.4	4.4	1640	9	US-10-106-698-499		499, App	
O	37	34.4	4.4	1665	2	US-09-919-002-8655		٠,٠	
O	38	34.4	4.4	5131	-	PCT-US02-09944-128		128, App	
O	39	34.2	4.4	646	9	US-10-027-632-210375		5	
	40	34.2	4.4	629	9	US-10-027-632-282527		282527,	
O	41		4.4	1090	9	-632		254462,	
U		34	4.3	344	9	US-10-027-632-268832	Sequence	268832,	
	43	34	4.3	508	9	US-10-027-632-90677	Sequence	90677, A	
O	44	34	4.3	638	9	US-10-027-632-45221	Sequence		
Ų		34	4 3	638	9	-027-632	Sequence	45222, A	
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						ALIGNMENTS			
RES	RESULT								
-Sn	09-053	US-09-053-375B-226	326				٠		
	equenc	Sequence 226, Applic	Applic	ation U	s/0	Sequence 226, Application US/09053375B			
٠	ADDI TO	ABBITCANT: Chanchik Alex	TALLON:	A 10 v					
	APPLICANT.	ANT.	aibilas	Ribilashvilli.	2	Robert			
	TITLE	OF INVE	NOLLN	Nuclei	ď	TITLE OF INVENTION: Nucleic Acid Arrays			
	FILE R	FILE REFERENCE: CLON-006	E: CLC	900-N					
	CURREN	IT APPLI	CATION	NUMBER	D .:	CURRENT APPLICATION NUMBER: US/09/053,375B			
••	CURREN	CURRENT FILING DATE:	NG DATE	1998	-08 - 31	-31			
	NUMBER	S OF SEC	O ID NO						
	SOFTWARE	ARE: Fas	FastSEQ for		OWS	Windows Version 4.0			
••	EO ID	SEQ ID NO 226				*			
	LENGTH:	гн: 1816 Эчэ							
••	TYPE: DNA	DNA		1					
:	ORGAN	ORGANISM: HOMO		sapiens					
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Best Local Similarity 84.9
Matches 667; Conservative
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             caaattgcggcacatgtcataagtgaggccagcagtaaaacaacatctgtgttacagtgg
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                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Chenchik, Alex
APPLICANT: Chenchik, Matvey
TITLE OF INVENTION: Hematology/Immunology Array
TITLE REFERENCE: CLON-006CIP15
CURRENT APPLICATION NUMBER: US/09/442,384B
CURRENT FILING DAFE: 1999-11-17
PRIOR APPLICATION NUMBER: 09/053,375
PRIOR APPLICATION NUMBER: 09/053,375
NUMBER OF SEQ ID NOS: 830
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 763.6; DB Pred. No. 9e-195; 0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                       Sequence 457, Application US/09442384B
GENERAL INFORMATION:
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98.2%;
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Best Local Similarity 98.2
Matches 772; Conservative
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CRGANISM: homo
US-09-442-384B-457
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US-09-442-384B-457
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                                                                 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAAACAACATCTGTGTTACAGTGG
                                                                                                                                                  CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
aacaaagaggagagaagaaagaaàacagctttgaaatgcaaaaaggtgatcagaatcct
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Pred. No. 7.8e-147;
); Mismatches 116;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Bibliashvilli, Robert TITLE OF INVENTION: Nucleic Acid Arrays FILE REFERENCE: CLON-006 CURRENT APPLICATION NUMBER: US/09/053,375B CURRENT FILING DATE: 1998-08-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 1543
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 821
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CCAGTTTGAAGGCTTTGTGAAGGATATAATGTTAAACAAAGAGGAGAGAAGAAAAA
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; ORGANISM: Homo sapiens
US-09-875-453A-9
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APPLICANT:
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                   AACAAAGAGAGACGAAGAAAAGAAAACAGCTTTGAÄATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                               CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
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                                                            TTACTGAACTGTGAGGAGTTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTÄ
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Pred. No. 5e-87;
0; Mismatches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Farr, Spencer B.
APPLICANT: Pickett, Gavin G.
APPLICANT: Pickett, Robin Elleen
APPLICANT: Dunn, II, Robert Thomas
TITLE OF INVENTION: CANINE TOXICITY GENES
FILE REFERENCE: 400742000200
CURRENT APPLICATION NUMBER: US/09/911,904
CURRENT FILING DATE: 2002-04-09
PRIOR APPLICATION NUMBER: US 60/220,057
PRIOR FILING DATE: 2000-07-21
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Best Local Similarity 83.0%;
Matches 424; Conservative
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GS-09-911-904-131
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                                                                                                                                GGCCAGCAGTAAAACAACATCTGTTACAGTGGGCTGAAAAAGGATACTACACCATGAG
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TITLE OF INVENTION: PROMOTERS FOR REGULATED GENE EXPRESSION
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Pred. No. 2.1e-26;
0; Mismatches 30;
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CURRENT APPLICATION NUMBER: US/09/875,453A
CURRENT FILING DATE: 2002-05-06
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SOFTWARE: FastSEQ for Windows Version 4.0
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Lim, Moon Young
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Thomas, Rita L.
Kongpachith, Ana
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Best Local Similarity 83.6
Matches ,153; Conservative
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FastSEQ for Windows
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ilarity 49.6%;
Conservative
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US-09-919-002-664
                                            TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                        Similarity
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LENGTH: 2127
                                                                            US-10-112-699-202
                             569
                                                                                                                                                           117;
             SEQ ID NO 202
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Best Local S:
Matches 117;
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NUMBER OF
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                                                              ATGAAGATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
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   ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC 60
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                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Gaiger, Alexander
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches 119; Indels
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Pred. No. 0.012;
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CURRENT APPLICATION NUMBER: PCT/US02/10421
CURRENT FILING DATE: 2002-03-28
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CURRENT APPLICATION NUMBER: US/10/112,699
CURRENT, FILING DATE: 2002-03-28
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cal Similarity 49.6%;
117; Conservation
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Wang, Siging
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; ORGANISM: Homo sapiens
PCT-USO2-10421-202
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LENGTH: 569
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                                                                                                                                    ttcatacattacagtggttggaataaaaattgggatgaatgggttccggagagcagagta 224
                                               Gaps
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PRIOR APPLICATION NUMBER: APPLICATION NUMBER: 09/359,922
PRIOR FILING DATE: FILING DATE: 1999-07-22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches 120;
                                            0; Mismatches 119;
    DB 6;
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  Score 45.6; DB Pred. No. 0.012;
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Pred. No. 0
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Best Local Similarity 49.2%;
Matches 116; Conservative
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JANT: Mullahy, Sara J.
JANT: Naughton, Rebecca E.
OF INVENTION: POLYNUCLECTIDES OF URINARY TRACT TISSUE
REFERENCE: PD-1037 CIP
                                                                                                                                                                                                  CATION NUMBER: US/09/540,210B
Application US/09540210B
                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER: 60/014,010
March 25, 1996
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ING DATE: July 31, 1996
LICATION NUMBER: 08/903,556
                                                                                                                                                                                                                               TION NUMBER: 08/972,899
                                                                                                                                                                                                                                                                                                                                 NG DATE: September 27, 1996
ICATION NUMBER: 60/005,526
                                                                                                                                                                                                                                                                                                                                                                         September 29, 1999
NUMBER: 08/824,029
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NUMBER: 08/862,178
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MIMBER: 60/018,217
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NUMBER: 08/881,589
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ICATION NUMBER: 60/023,308
NG DATE: July 31, 1996
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FILING DATE: August 1, 1997
APPLICATION NUMBER: 60/025,204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PPLICATION NUMBER: 60/025,217
ILING DATE: August 22, 1996
PPLICATION NUMBER: 08/937,142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08/903,802
                                                                                                                                                                                                                                                      November 18, 1997
UMBER: 08/395,244
February 27, 1995
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ATION NUMBER: 08/903,471
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PPLICATION NUMBER: 60/026,598
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DATE: July 30, 1997
                                     Seilhamer, Jeffrey J.
Delegeane, Angelo M.
Stuart, Susan G.
Stuve, Laura L.
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323 AAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCTCAAATTGCGGCACATGTCATAA 382
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   121 gggctgccccaggaaagaagacatctggtctgcaacagaaaaatgttgaagtgaaaacga 180
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Pred. No. 0.41;
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NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No: hu00141818
LOCATION: 24, 113
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: a, t, c, g, or other US-09-540-2108-7000
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GENERAL INFORMATION:
APPLICANT: Sellhamer, Jeffrey J.
APPLICANT: Delegene, Angelo M.
APPLICANT: Stuart, Susan G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: May 29, 1997
APPLICATION NUMBER: 09/107,592
FILING DATE: June 30, 1998
PEPLICATION NUMBER: 60/052,751
FILING DATE: July 1, 1997
APPLICATION NUMBER: 09/094,079
                                                                                                                                                                                                                                  PELICATION NUMBER: 60/040,431
TILING DATE: MARCH 5, 1997
APPLICATION NUMBER: 09/041,894
FILING DATE: MARCH 12, 1998
APPLICATION NUMBER: 60/040,199
                                                                                                                                                                                                                                                                                                                                                                                                                               ILING DATE: March 14, 1997
PPLICATION NUMBER: 09/050,817
ILING DATE: March 30, 1998
PPLICATION NUMBER: 60/043,792
ILING DATE: April 11, 1997
                                                                                                                           ILING DATE: February 13, 1997
PPLICATION NUMBER: 09/035,172
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PPLICATION NUMBER: 60/048,431
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Best Local Similarity 48.8
Matches 103; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Homo sapiens
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APPLICANT: Stuve, Laura L.
APPLICANT: Mullahy, Sara J.
APPLICANT: Mullahy, Saca J.
TITLE OF INVENTION: POLYNUCLEOTIDES OF URINARY TRACT TISSUE
                                                                                    CURRENT APPLICATION NUMBER: US/09/540,210B CURRENT FILING DATE: 2002-04-03
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ING DATE: August 22, 1996
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R: 08/881,589
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TON NUMBER: 08/905,881
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                                                                                                                   PRIOR APPLICATION NUMBER: 08/972,899
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APPLICATION NUMBER: 60/030,14
                                                                                                                                    FILING DATE: November 18, 1997
APPLICATION NUMBER: 08/395,244
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                                                                                                                                                                                                    FILING DATE: September
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298 TTAAACAAAGGGGGGGGAGGAAAGAAAGGCTTTGAAATGCAAAAAGGTGATCAGAAT 357
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Wang, David G. Application and Mapping of Single Nucleotide TITLE OF INVENTION: Identification and Mapping of Single Nucleotide TITLE OF INVENTION: Polymorphisms in the Human Genome FILLE REFERENCE: 10827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILLING DATE: 2002-04-30
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Pred. No. 0.85;
0; Mismatches · 88;
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; OTHER INFORMATION: Incyte ID No: hu00678829
US-09-540-210B-14649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , Application US/10027632
                                                                                                                                                                                                                                                             APPLICATION NUMBER: 60/052,751 FILING DATE: July 1, 1997 APPLICATION NUMBER: 09/094,079
APPLICATION NUMBER: 09/041,894
FILING DATE: March 12, 1998
APPLICATION NUMBER: 60/040,199
                                                                                                          APPLICATION NUMBER: 60/043,792
FILING DATE: April 11, 1997
APPLICATION NUMBER: 09/074,999
                                                 FILING DATE: March 14, 1997
APPLICATION NUMBER: 09/050,817
                                                                                                                                                                       FILING DATE: MAY 8, 1998
APPLICATION NUMBER: 60/048,431
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Best Local Similarity 50.8%;
Matches 91; Conservative
                                                                                       FILING DATE: March 30,
                                                                                                                                                                                                                                                     June 30,
                                                                                                                                                                                                              FILING DATE: May 29,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Homo sapiens
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GENERAL INFORMATION:
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LING DATE: 2000-02-24
DPLICATION NUMBER: US 60/167,363

FILING DATE: 2000-03-29 APPLICATION NUMBER: US 60/185,218

PRIOR PRIOR RIOR

PRIOR APPLICATION NUMBER: US 60/218,006

2000-07-12 2000-04-20

PRIOR FILING DATE: 2000-07 PRIOR APPLICATION NUMBER:

PRIOR FILING DATE: 2000-0-PRIOR APPLICATION NUMBER:

US 60/198,676 US 60/193,483

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NUMBER OF SEQ ID NOS: 325720
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Best Local Similarity 51.55
Matches 88; Conservative
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                                                                                                                        US-10-027-632-297617
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US-10-027-632-297617
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                                                                                                                                                                                                                                                                                                           GAGCATGAAGATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTC 116
                                                                                                                                                                                                                                                                                                                                                                                                                               250 aatattaagtactttttattggctaagtccagtacaaatttagaaacagcaatgtatca 309
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                                                                                                                                                                                                                                                                                                                                                 190 gagattagtgattattattcactatttggcagccttactttttcagacattttgatgcc 249
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                                                                                                                                                                                                                           Length 620;
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                                                                                                                                                                                                                                                                   83; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Sing
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       83;
                                                                                                                                                                                                                           DB 6;
                                                                                                                                                                                                                       Score 38.2; DB Pred. No. 1.2; 0; Mismatches
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51.5%; Pred. No. 1.2
                                   NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 35707
LENGTH: 620
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PRIOR FILING DATE: 2000-07-12
PRIOR PLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR PLICATION UNMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 60/167,363 FILING DATE: 1999-11-23 APPLICATION NUMBER: US 60/156,358 FILING DATE: 1999-09-28
PRIOR APPLICATION NUMBER: US 60/146,002 PRIOR FILING DATE: 1999-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US 60/146,002 PRIOR FILING DATE: 1999-08-09
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 62305, Application US/10027632
                                                                                                                                                                                                                         cch 4.9%;
al Similarity 51.5%;
88; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     88; Conservative
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Best Local Similarity
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Matches 88; Conserv
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                                                                                                                                      ORGANISM: Human
US-10-027-632-35707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-027-632-62305
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TYPE: DNA
; ORGANISM: Human
US-10-027-632-62305
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SEQ ID NO 62305
                                                                                                                        TYPE: DNA
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Sequence 206146, Application US/10027632
GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Dolymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT PAPLICATION WOMBER: US/10/027,632
CURRENT FILING DATE: 2002-04.30
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR PAPLICATION NUMBER: US 60/198,676
PRIOR APPLICATION NUMBER: US 60/198,676
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              57 GAGCATGAAGATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTC 116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     190 gagattagtgattattattcactatttggcagccttactttttcagacattttgatgcc 249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          117 AGCACTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGGAATGT 176
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      310 tcgtgtagatgttcatttcatttaaatgaatggcactatgaacaaagtaca 360
177 TCATGAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGA 227
                       DB 6; Length 620;
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Pred. No. 1.2;
0; Mismatches
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APPLICATION NUMBER: US 60/193,483
FILING DATE: 2000-03-29
APPLICATION NUMBER: US 60/185,218
FILING DATE: 2000-02-24
                                                                                                                                 Sequence 297617, Application US/10027632 GENERAL INFORMATION:
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51.58;
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us-08-982-272-6.rnpn

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236 TATCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAA 295
                                                                                                                                                                                                                                                                                                                                                                                                                                                            176 TTCATGAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCT 235
                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Wang, David G. TITLE OF INVENTION: Identification and Mapping of Single Nucleotide TITLE OF INVENTION: Polymorphisms in the Human Genome
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  296 IGTTAAACAAAGAGGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTG 349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      gagatatcagggagctgaaaatacaatgaataatgtgaaaatgcaatagagag 346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 6; Length 663;
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4.8%; Score 38; DB 6

Best Local Similarity 51.1%; Pred: No. 1.3;
Matches 89; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                     Score 38; DB Pred. No. 1.3;
                                                                                                                                                                                                                                                                                                                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
                                                                                                           NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 206146
LENGTH: 663
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PLICATION NUMBER: US 60/185, 218
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LICATION NUMBER: US 60/156,358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATE: 2000-07-12
ION NUMBER: US 60/198,676
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APPLICATION NUMBER: US 60/193,483
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LICATION NUMBER: US 60/167/363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , Application US/10027632
                                                                     PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
APPLICATION NUMBER: US 60/167,363
                   TLING DATE: 1999-11-23
PPLICATION NUMBER: US 60/156,358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LE REFERENCE: 108827,129
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Best Local Similarity 51.1%;
Matches 89; Conservative
                                                       ILING DATE: 1999-09-28
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US-10-027-632-206147
                                                                                                                                                                                               TYPE: DNA
ORGANISM: Human
JS-10-027-632-206146
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1,176 TTCATGAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAAAAGATCCT 235 ttaaagaaaactcagtgaacatcaagaaaatgcaatgaaacaattcaacaaaatgagaaa 232

Indels

85;

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236 TATCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAA 295
             296 TGTTAAACAAAGGAGGAGGAAGAAAGAAAAAAAAAAGCTTTGAAATGCAAAAAGGTG 349
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Search completed: May 30, 2002, 05:40:03 Job time: 16777 sec

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May 30, 2002, 02:51:38 ; Search time 139.75 Seconds (without alignments) 1381.524 Million cell updates/sec
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786
1 ATGATAGAAACATACAGCCA.....TTGGCTTACTCAAACTCTGA 786
GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

/cgn2_6/ptodata/2/ina/PCTUS_COMB.seq /cgn2_6/ptodata/2/ina/backfiles1.seg

/cgn2_6/ptodata/2/ina/5A_COMB.seq /cgn2_6/ptodata/2/ina/5B_COMB.seq /cgn2_6/ptodata/2/ina/6A_COMB.seq

Issued_Patents_NA:

Database :

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Minimum DB seq length: 0 Maximum DB seq length: 200000000

SUMMARIES

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US-08-446-922-3	PCT-US93-10034-3 US-07-940-605A-1	US-08-184-422-7	US-08-360-923A-1	US-08-431-055-3	US-08-690-096-1	US-08-249-189-11	US-08-484-624A-11	US-08-477-733B-11	US-08-763-995-1	US-09-088-913A-11	US-08-589-771B-7	US-08-769-819-11	US-08-770-974-11	US-08-858-197-3	US-08-249-189-15	US-08-484-624A-15	US-08-477-733B-15	US-09-088-913A-15	US-08-769-819-15	US-08-770-974-15	US-08-446-922-10	US-08-249-189-20	US-08-484-624A-20	US-08-477-733B-20	US-09-088-913A-20
786 1	786 5 840 1	840 1	840 1	840 1	840 2	840. 2	840 2	840 2	840. 3	840 3	840 3	840 4	840 4	840 4	425 2	425 2	425 2	.425 3	425 4	425 4	929 1	929 2	929 2	929 2	929 3
97.2	97.2	97.2	97.2	97.2	97.2	97.2	97.2	97.2	97.2	97.2	97.2	97.2	97.2	97.2	81.2	81.2	81.2	81.2	7.					81.1	81.1
763.6	763.6	763.6	763.6	763.6	763.6	763.6	763.6	763.6	763.6	763.6	763.6	763.6	763.6	763.6	638.2	638.2	638.2	638.2	638.2	638.2	637.2	637.2	637.2	637.2	637.2
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	.6 97.2 786 1 US-08-446-922-3 Sequence 3,	.6 97.2 786 1 US-08-446-922-3 Sequence 3, 6 97.2 786 5 PCT-US93-10034-3 Sequence 3, 6 97.2 840 1 IIS-07-940-6058-1	.6 97.2 786 1 US-08-446-922-3 Sequence 3, 6 97.2 786 5 PCT-US93-10034-3 Sequence 3, 6 97.2 840 1 US-08-184-422-7 Sequence 1, 6 97.2 840 1 US-08-184-422-7	.6 97.2 786 1 US-08-446-922-3 Sequence 3, 6 97.2 786 5 PCT-US93-10034-3 Sequence 3, 6 97.2 840 1 US-08-184-422-7 Sequence 7, 6 97.2 840 1 US-08-184-422-7 Sequence 7, 6 97.2 840 1 US-08-1860-923A-1	6 97.2 786 1 US-08-446-922-3 Sequence 3, 6 97.2 786 5 PCT-US93-10034-3 Sequence 3, 6 97.2 840 1 US-08-940-655-1 Sequence 1, 6 97.2 840 1 US-08-184-422-7 Sequence 7, 6 97.2 840 1 US-08-340-923A-1 Sequence 3, 6 97.2 840 1 US-08-431-055-3 Sequence 3,	6 97.2 786 1 US-08-446-922-3 Sequence 3, P. 786 5 PCT-US93-10034-3 Sequence 3, P. 6 97.2 840 1 US-07-940-605A-1 Sequence 1, P. 6 97.2 840 1 US-08-184-422-7 Sequence 7, P. 6 97.2 840 1 US-08-340-923A-1 Sequence 1, P. 6 97.2 840 2 US-08-690-096-1 Sequence 3, P. 6 97.2 840 2 US-08-690-096-1	6 97.2 786 1 US-08-446-922-3 Sequence 3, 8 6 97.2 786 5 PCT-US93-10034-3 Sequence 3, 8 6 97.2 840 1 US-07-940-605A-1 Sequence 1, 8 6 97.2 840 1 US-08-184 422-7 Sequence 7, 8 6 97.2 840 1 US-08-360-923A-1 Sequence 1, 8 6 97.2 840 1 US-08-341-055-3 Sequence 1, 8 6 97.2 840 2 US-08-690-096-1 Sequence 1, 9 840 2 US-08-690-096-1 Sequence 1, 9 840 2 US-08-249-189-11 Sequence 1, 9 80 2 US-08-249-189-11 Sequence 1, 9 80 2 US-08-249-189-11	6 97.2 786 1 US-08-446-922-3 Sequence 3, A 6 97.2 786 5 PCT-US93-10034-3 Sequence 3, A 6 97.2 840 1 US-08-446-922-7 Sequence 1, A 6 97.2 840 1 US-08-1444-422-7 Sequence 7, A 6 97.2 840 1 US-08-360-938-1 Sequence 1, A 6 97.2 840 1 US-08-431-055-3 Sequence 1, A 8 840 2 US-08-491-18-11 Sequence 1, A 6 97.2 840 2 US-08-249-189-11 Sequence 11, A 6 97.2 840 2 US-08-444-624A-11 Sequence 11, A 6 97.2 840 2 US-08-444-624A-11 Sequence 11, A 6 97.2 840 2 US-08-448-624A-11	6 97.2 786 1 US-08-446-922-3 Sequence 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Sequence 3, p. 6 97.2 840 1 US-08-446-922-7 Sequence 1, p. 6 97.2 840 1 US-08-360-954-1 Sequence 7, p. 6 97.2 840 1 US-08-360-933A-1 Sequence 1, p. 6 97.2 840 1 US-08-360-951 Sequence 1, p. 6 97.2 840 2 US-08-249-189-11 Sequence 1, p. 6 97.2 840 2 US-08-249-189-11 Sequence 11, c. 6 97.2 840 2 US-08-484-624A-11 Sequence 11, p. 6 97.2 840 3 US-08-95-1 Sequence 11, p. 6 97.2 840 3 US-08-95-1 Sequence 11, p. 6 97.2 840 3 US-08-95-1 Sequence 11, p. 6 97.2 840 3 US-08-96-1 Sequence 11, p. 6 97.2 840 3 US-08-913A-11 Sequence 11, p. 6 97.2 840 3 US-08-96-1 Sequence 11, p. 6 97.2 840 3 US-08-913A-11	6 97.2 786 5 PCT-US93-10034-3 Sequence 3, p. 6 97.2 786 5 PCT-US93-10034-3 Sequence 3, p. 6 97.2 840 1 US-08-446-922-7 Sequence 1, p. 6 97.2 840 1 US-08-144 4.22-7 Sequence 7, p. 6 97.2 840 1 US-08-340-9233-1 Sequence 1, p. 6 97.2 840 1 US-08-491-1 Sequence 1, p. 6 97.2 840 2 US-08-491-1 Sequence 1, p. 6 97.2 840 2 US-08-491-1 Sequence 11, c. 6 97.2 840 2 US-08-491-1 Sequence 11, c. 6 97.2 840 3 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Sequence 1, 763.6 97.2 84.0 2 US-08-809-90-1 Sequence 1, 763.6 97.2 84.0 2 US-08-849-189-11 Sequence 11, 763.6 97.2 84.0 2 US-08-484-624A-11 Sequence 11, 763.6 97.2 84.0 3 US-08-913A-11 Sequence 17, 763.6 97.2 84.0 4 US-08-913A-11 Sequence 7, 763.6 97.2 84.0 4 US-08-913A-11 Sequence 7, 763.6 97.2 84.0 4 US-08-79-11 Sequence 11, 763.6 97.2 84.0 4 US-08-79-11	763.6 97.2 786 1 US-08-446-922-3 Sequence 3, 763.6 97.2 786 5 PCT-US93-10034-3 Sequence 3, 763.6 97.2 786 5 PCT-US93-10034-3 Sequence 1, 763.6 97.2 84.0 1 US-08-144-422-7 Sequence 7, 763.6 97.2 84.0 1 US-08-130-131-055-3 Sequence 1, 763.6 97.2 84.0 1 US-08-431-055-3 Sequence 1, 763.6 97.2 84.0 2 US-08-690-096-1 Sequence 1, 763.6 97.2 84.0 2 US-08-690-196-11 Sequence 11, 763.6 97.2 84.0 2 US-08-691-199-11 Sequence 11, 763.6 97.2 84.0 2 US-08-691-199-11 Sequence 11, 763.6 97.2 84.0 3 US-09-088-913A-11 Sequence 11, 763.6 97.2 84.0 3 US-09-088-913A-11 Sequence 11, 763.6 97.2 84.0 4 US-08-769-191-1 Sequence 11, 763.6 97.2 84.0 4 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Sequence 20, Appl	20,	Sequence 5, Appli	Sequence 1, Appli	Sequence 5, Appli	1,	1, 7	Sequence 22, Appl											
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637.2	637.2	584.4	584.4	584.4	584.4	584.4	. 584.4	584.4	584.4	582.8	582.8	445.6	445.6	445.6	445.6	445.6	445.6	
28	29	30	31	32	33	34.	35	36	37	38	39	40	41	42	43	44	45	

ALIGNMENTS

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Springs, Melanie
: Srinivasan, Subhashini .
INVENTION: Methods of Preparing Soluble, Oligomeric
INVENTION: Proteins
                                                                                                                                                                                                                                                                                                         Word for Apple, Version 5.1a
                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: Apple Operating System 7.1 SOFTWARE: Microsoft Word for Apple, Version CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                   STRET: 51 University Street
CITY: Seatile
STATE: WA
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                       US/08/446,922
se 3, Application US/08446922
No. 5716805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,69
                                                                                                                                                                                                                                                                         Apple Macintosh
                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   : 786 base pairs
nucleic acid
                                                                                                            NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Immunex C
                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER:
                                                                              TITLE OF INVENTION:
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                               GENERAL INFORMATION:
APPLICANT: Sprigg
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Hum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: HYPOTHETICAL:
                                                                                                                                                                                                                       98101
                                                                                                                                                                                                                                                                         COMPUTER:
                                                                  APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 540
                                                                                                                                                                                                                                                                                                                                                                                                                                     541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 600
                                                                                                                                                                                                                                                                              GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG 480
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                                                                                                                                                          CITITIGCIGIGIAICTICAIAGAAGGIIGGACAAGAIAGAAGAIGAAAGGAAICIICAI 180
                                                                                                                                                                                                                      240
                                                                                                                               1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC 60
                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                           0
                                           Length 786;
                                                                Indels
                                              Score 763.6; DB 1;
Pred. No. 1.3e-213;
0; Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL SPINION:
APPLICANT: STINIVASAN, SUBMASHINI
TITLE OF INVENTION: Methods of Preparing
                                               97.2%;
                                                                   Conservative
                                                            Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   111111
CTCTGA 786
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PCT-US93-10034-3
; NAME/KEY:
; LOCATION:
US-08-446-922-3
                                                                     772;
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Best Local S:
Matches 772
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241 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 786;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 763.6; DB 5;
Pred. No. 1.3e-213;
0; Mismatches 14;
                                                                                                                                                                  Version #1.25
                                                                                                                           MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, V
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/10034
                                   E: Immunex Corporation
51 University Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0
                                                                                                                                                                                                                                                 NAME: Perkins, Patricia A REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    97.28;
                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 98.2
Matches 772; Conservative
                                                                                                                                                                                                                                                                                                                                                          786 base pairs
nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                     single
                                                                                                                 COMPUTER READABLE FORM:
                           CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                    linear
           SEQUENCES
FITTE OF INVENTION:
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                                                                                                                                                                                                                         CLASSIFICATION:
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                                                 STREET: 51 vm.
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                                                                                          USA
                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE:
                                                                                                                                                                                                           FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION:
PCT-US93-10034-3
                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY:
                                       ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                          LENGTH:
                                                                                            COUNTRY:
               NUMBER OF
                                                                               STATE:
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                                                                                1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGGGAGC 60
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      Length 840;
                                            Indels
                                              14;
    Score 763.6; DB 1;
Pred. No. 1.3e-213;
                                        0; Mismatches
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Patent No. 556531
GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
APPLICANT: DAVISON, BARRY
97.2%;
                                            Conservative
                       Similarity
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GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780
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                                                                                                                                                                                                                                          CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT 540
                                                                                                                                                                                                                                                             301 AACAAAGAGGAGACGAAGAAAGAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 360
                                                                                                             361 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 420
                                                                                                                              GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAAAG 480
                                                                                                                                                                                            601 TTCGAGAGAATCTTACTCAGAGGTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                                                                                                                                                                           CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                                               DISEASE CHARACTERIZED BY NEOPLASTIC CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh Apple Macintosh System 7.1
OPERATING SYSTEM: Apple Macintosh, Version #5.1a
SOFTWARE: Microsoft Word for Macintosh, Version #5.1a
CURRENT APPLICATION DATE:
FILING DATE: December 21, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: METHOD OF PREVENTING OR TREATING TITLE OF INVENTION: DISEASE CHARACTERIZED BY NEOPLAST TITLE OF INVENTION: EXPRESSING CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USSN 08/172,664
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application US/08360923A Patent No. 5674492
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             December 23,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MURPHY, WILLIAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LONGO, DAN L
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX: C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: SEATTLE STATE: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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APPLICANT:
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                                                                             TITLE OF INVENTION: DETECTION AND TREATMENT OF MUTATIONS TITLE OF INVENTION: IN A CD40 LIGAND GENE NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 763.6; DB 1;
Pred. No. 1.3e-213;
0; Mismatches 14;
                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: Apple Operating System 7.1 SOFTWARE: MS Word for Apple 5.1, Version a CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/08/184,422
                                                                                                                                                                                                                                                                                                                                                                                                 PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/009,258
FILING DATE: 01/22/93
                                                                                                                                                     : IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: PERKINS, PATRICIA ANNE REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 2810 TELECOMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        97.2%;
98.2%;
                                                                                                                                                                                                                                                                              COMPUTER: Apple Macintosh
                                                        SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 01/22/93 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                        Floppy disk
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Best Local Similarity 98.2
Matches 772; Conservative
                                                                        WIDMER, MICHAEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 840 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              single
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 800
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                        FANSLOW,
RENSHAW,
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LOCATION:
US-08-184-422-7
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CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
                                                                                                61 ATGAAGATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
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                                1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC
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                                                                               GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAA
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                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: KEHRY, MERILYN R
APPLICANT: CASTLE, BRIAN E
TITLE OF INVENTION: METHODS FOR PROLIFERATING AND
TITLE OF INVENTION: DIFFERENTIATING B CELLS, AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
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Pred. No. 1.3e-213;
); Mismatches 14;
                                                                                                                                                                                                                                                                                                                                                                                                3: STERN, KESSLER, GOLDSTEIN & FOX 100 NEW YORK AVE. N.W. SUITE 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floopy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver
APPLICATION NUMBER: US/08/431,055
FILING DATE: 28-APR-1995
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US 08/234,580 FILING DATE: 28-APR-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                         Sequence 3, Application US/08431055 Patent No. 5817516
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                36,217
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 97.2%;
Best Local Similarity 98.2%;
Matches 772; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (202)371-2600
TELEFAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: MILLMAN, ROBERT A REGISTRATION NUMBER: 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
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STATE: D.C.
77. 20005
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; LOCATION:
US-08-431-055-3
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Pred. No. 1.3e-213;
0; Mismatches 14;
           NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 28 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                               97.2%;
                                                                                                           INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                                                                            TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
ATTORNEY/AGENT INFORMATION
                                                                                                                                       LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 97.2
Best Local Similarity 98.2
Matches 772; Conservative
                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                                                                                                  NAME/KEY:
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US-08-360-923A-1
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361 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACAACATCTGTGTTACAGTGG 420
                                             240
                                                                                      241 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
                                                                                                  AACAAAGAGAGAGAGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 360
                                                                                                                                            GCTGAAAAAGGATACTACACCATGAGCAACATTGGTAACCCTGGAAAATGGGAAACAG
                                                                                                                                                                                                                                                                         GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC
                                                         PatentIn Release #1.0, Version #1.25
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HOLLENBAUGH, DIANE
LEDBETTER, JEFFREY A.
NYENTION: SOLUBLE LICANDS FOR CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOPTWARDER: NEATHER.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/08690096
Patent No. 5945513
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New York
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APPLICANT:
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US-08-690-096-1
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                                                                                                                                                                                                                                                                                                                                                 Score 763.6; DB 2;
Pred. No. 1,3e-213;
); Mismatches 14;
                                                      APPLICATION NUMBER: US 07/940,605 FILING DATE: 04-SEP-1992 ATTORNEY/AGENT INFORMATION:
JMBER: US/08/690,096
                                                                                                            18,872
                                                                                                                                   TELECOMMUNICATION INFORMATION:
TELEPHONE: 212 790-9090
                                                                                                                                                            TELEFAX: 212 869 8864/9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                  DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                       97.2%;
                                                                                                Misrock, S. Leslie
                                                                                                                        REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                               LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 98.2 ges 772; Conservative
                                              PRIOR APPLICATION DATA:
                                                                                                     NAME: Misrock, S. Le
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                         CDS 22..807
                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                   Similarity.
                   FILING DATE: 3. CLASSIFICATION:
                                                                                                                                                                                                                                                                  MOLECULE TYPE:
                                                                                                                                                                                                                                                                                         , NAME/KEY:
LOCATION:
US-08-690-096-1
                                                                                                                                                                                                                                                        ropology:
                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                                                                                                                                                                                                                                              FEATURE
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Matches
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180

360

381

321

300

240

261

420 441 480 ; 0

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IMMEDIATE SOURCE:
CLONE: CD40-L
                                                   NAME/KEY:
                 CLONE:
                                FEATURE
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                                 661 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
                                                                                                                  622 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 681
                                                  GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Operating System 7.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
COMPUTER: Apple Macintosh
SOFTWARE: Microsoft Word for Apple,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 23
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US/08/249,189
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
CLASSIFICATION 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AUDRESSEE: IMMUNEX CORPORATION STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                       Sequence 11, Application US/08249189 Patent No. 5961974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 28
FELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WASHINGTON: USA
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MOLECULE TYPE: cDNA
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                                                                                                                                                                  781 CTCTGA 786
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802 CTCTGA 807
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                                                                                                                                                                                                                                                      RESULT 8
US-08-249-189-11
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                                                                                                                                                                                                                                                                             ATGAAGATTTTTATGTATTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGC 120
                                                                                                                                                                                                                                                                                                                                                                            CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
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                                                                                                                                                                                                                                                                                                                              ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGC
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                                                                                                                                     Gaps
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                                                                                         840;
                                                                                         Length
                                                                                                                                     Indels
                                                                                     .6; DB 2;
1.3e-213;
                                                                                                                                     Mismatches
                                                                                       Score 763.6;
Pred. No. 1.3
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US-08-484-624A-11
Sequence 11, Application US/08484624A
; Patent No. 5962406
                                                                                  97.2%;
illarity 98.2%;
Conservative
46..831
                                                                                       Query Match
Best Local Similarity
Matches 772; Conserv
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; LOCATION:
US-08-249-189-11
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661 CAATCCATTCACTTGGGAGGAGTATTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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                                                                                                             CTTTTTGCTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
                                               ATGAAGATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
                                                              601 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
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                                                                                                                               181 GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC
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APPLICANT: GTBSON, MARYLOU
APPLICANT: MORRIS, ARVIA E.
APPLICANT: MORSEW, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 26
CORRESPONDENCE ADDRESS:
ANDRESS ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 11, Application US/08477733B Patent No. 5981724 GENERAL INFORMATION:
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826 CTCTGA 831
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APPLICANT:
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pred. NO. 1.3e-213;
); Mismatches 14; Indels 0;
                                                                                                                                                                                                                                                                                                                                 COMPUTER: Apple Macintosh OPERATING SYSTEM 7.5.5 OPERATING SYSTEM: Apple Operating System 7.5.5 SCHWARE: MS WORD for Apple Power Macintosh, version 6.0.1 SOFTWARE: APPLICATION DATA: APPLICATION DATA:
                                                                                               APPLICANT: MORRIS, ARVIA E.
APPLICANT: MCGREW, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRICR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
                                                                                       SRINIVASAN, SUBHASHINI
GIBSON, MARYLOU
                                                                                                                                                                                                   SEE: IMMUNEX CORPORATION : 51 UNIVERSITY STREET SEATTLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     08/477,733
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 07/969,70:
FILING DATE: October 23, 1992
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 280
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,
FILING DATE: June 07, 1995
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                                           ARMITAGE, RICHARD
                                                           PANSLOW, WILLIAM SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 98.2
Matches 772; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTIČS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
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                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
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ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-484-624A-11
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                                                                                                                                                                                                                                                                                          98101
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                                                                                                                                                                                                        ADDRESSEE:
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                                                                                                                                                                                                                                                                         COUNTRY:
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IMMUNEX CORPORATION 51 UNIVERSITY STREET

ADDRESSEE

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361 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 600
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                                                               301 AACAAAGAGAGACGAAGAAAAGAAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                481 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                       241 TTACTGAACTGTGAGGAGTTAAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: MARRASKOVSKY, EUGENE
TITLE OF INVENTION: METHOD OF ACTIVATING DENDRITIC CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Macintosh, Version #6.0.1
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/763,995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/677,762
FILING DATE: 10 JUL 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: Power Macintosh 7200/90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TITLE OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/08763995
Patent No. 6017527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
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STATE: WASHING?
COUNTRY: USA
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| 826 CTCTGA 831
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SOFTWARE: MS Word for Apple Power Macintosh, version 6.0.1
CURRENT APPLICATION DATA:
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Pred. No. 1.3e-213;
0; Mismatches 14;
                                                                                                                                                                                                      APPLICATION NUMBER: US/08/477,733B
FILING DATE: June 07, 1995
                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/249,189
FILING DATE: MAY 24, 1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION WIBER: 07/969,703
FILING DATE: OCCODER 23, 1992
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
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APPLICATION NUMBER: 07/783,707
FILING DATE: 0ctober 25, 1991
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
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                                                                                                                        COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple O
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ONIGINAL SOURCE:
ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity 98.2
Matches 772; Conservative
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2065870606
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                 WASHINGTON
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US-08-477-733B-11
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                                                                    766 GIGACIGAICCAAGCCAAGIGAGCCAIGGCACIGGCTICACGICCIIIGGCIIACICAAA 825
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                                                                                                                                                                                                                                                                                                                                         NOVEL CYTOKINE THAT BINDS CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Apple Operating System 7.5.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/088,913A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 07/805,723 FILING DATE: December 5, 1991 PRIOR APPLICATION DATA: APPLICATION NUMBER: 07/783,707 FILING DATE: October 25, 1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 08/477,733
FILING DATE: June 07, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
                                                                                                                                                                          ESULT 12
5-09-088-9134-11
Sequence 11, Application US/09088913A
Patent No. 6087329
                                                                                                                                                                                                                                                                                        SRINIVASAN, SUBHASHINI
GIBSON, MARYLOU
MORRIS, ARVIA E.
MCGREW, JEFFERY
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APPLICATION NUMBER: 08/484,624
FILING DATE:
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51 UNIVERSITY STREET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER: 28C
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OPERATING SYSTEM: Apple O
                                                                                                                                                                                                                                                             FANSLOW, WILLIAM SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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nucleic acid
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WASHINGTON
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                                                                                                                                                                                                                                 GENERAL INFORMATION:
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ORIGINAL SOURCE:
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| 826 CTCTGA 831
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                      REGISTRATION NUMBER: 34,693
REPERENCE/DOCKET UNMBER: 28,
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                          Query Match 97.2%;
Best Local Similarity 98.2%;
Matches 772; Conservative
                                                                                          INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                              (206)233-0644
                                                                                                                                                                                                                                     ORGANISM: Homo sapiens IMMEDIATE SOURCE:
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LOCATION:
US-08-763-995-1
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                                                                                                                                    Gaps
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                                                                                                          Length
                                                                                                                                    Indels
                                                                                                          DB 3;
                                                                                                         Score 763.6; DB 3;
Pred. No. 1.3e-213;
                                                                                                                                   0; Mismatches
                                                                                                          97.28;
ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
                                                                                                                                   Matches 772; Conservative
                                             CDS
46..831
                                                                                                          Query Match
Best Local Similarity
                      CD40-L
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                                             ; NAME/KEY:
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US-09-088-913A-11
                      CLONE:
                                   FEATURE
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46 ATGATCGAAACATACAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 105
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                                                                                                       TREATMENT OF INDIVIDUALS EXHIBITING DEFECTIVE CD40L (as amended)
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Pred. No. 1.3e-213;
0; Mismatches 14;
                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC Compatible
OPERATING SYSTEM: MS-DOS/Windows 95
SOFTWARE: Word for Windows 95, 7.0a
                                                                                                                                                                                                                                                                                                                                                                                                      US/08/589,771B
                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/589,777
FILING DATE: January 22, 1996
CLASSIFICATION A35
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/009,258
FILING DATE: 01/22/93
                                                                                                                                                                             E: IMMUNEX CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: HENRY, JANIS C.
REGISTRATION NUMBER: 34,347
REFERENCE/DOCKET NUMBER: 2810
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                   WILLIAM
BLAIR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 97.2%;
Best Local Similarity 98.2%;
Matches 772; Conservative (
                                                                   APPLICANT: SPRIGGS, MELANIE APPLICANT: WIDMER, MICHAEL
                                                                                                                                                                                                                                                                                                               Floppy disk
                  DAVISON, BARRY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                           NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                      COUNTRY: 1157
                                   FANSLOW,
RENSHAW,
                                                                                                       TITLE OF INVENTION:
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ORGANISM: Hom
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US-08-589-771B-7
APPLICANT:
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                                 APPLICANT:
APPLICANT:
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FEATURE:
                  APPLICANT
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; Sequence 7, Application US/08589771B

RESULT 13 US-08-589-771B-7 us-08-982-272-6.rni

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361 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 420
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APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
MUMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SYSTEM: Apple Operating System 7.1 Microsoft Word for Apple, version
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               sequence 11, Application US/08769819
Patent No. 6264951
GENERAL INFORMATION:
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51 UNIVERSITY STREET
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FANSLOW, WILLIAM
SPRIGGS, MELANIE
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MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 'OR
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STATE: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  781 CTCTGA 786
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US-08-769-819-11
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APPLICANT:
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361. CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 420
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Pred. No. 1.3e-213
0; Mismatches 14
                                                                                                                                                                             NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 2802-E TELECOMMUNICATION INFORMATION:
         UMBER: 08/249,189
May 24, 1994
                                                    UMBER: 07/969,703
October 23, 1992
                                                                                                JMBER: 07/805,723
December 5, 1991
                                                                                                                                         07/783,707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        97.2%;
                                                                                                                                                     FILING DATE: October 25, ATTORNEY/AGENT INFORMATION:
07-JUN-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                840 base pairs
                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                    2065870430
                                                                                                           FILING DATE: December PRIOR APPLICATION NUMBER: 0
                                                                                                                                                                                                                                                                                                                             STRANDEDNESS: single
                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: (
                                                                                                                                                                                                                                                     2065870606
                                                                                                                                                                                                                                                                                                             nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: CDS
LOCATION: 46..831
               APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
nes 772; Conserv
                                                                                                                                                                                                                                                                                                                                           linear
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ORGANISM: HOMO
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                               FILING DATE:
                                                                        FILING DATE:
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Best Local S
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765
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                                                                                                                                                                                                                                                                                     CAATCCATTCACTTGGGAGGAGTATTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAT
                                                                                                 541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                                                                                                                                          TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                                                                                          706 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                          CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                     526 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintons
COMPUTER: Apple Macintons
COMPUTER: Apple Macintons
SOFTWARE: Microsoft Word for Apple, version 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/770,974
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: ARMITAGE, RICHARD
APPLICANT: SPRIGGS, MELLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVAGAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TILE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
CORRESPONDENCES: 24
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,733
FILING DATE: 02-AUG-1995
APPLICATION NUMBER: 08/249,189
FILING DATE: May 24, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: IMMUNEX CORPORATION STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 11, Application US/08770974
Patent No. 6290972
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            December 5, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
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FRY: USA
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1.3e-213;
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REFERENCE/DOCKET NUMBER: 2802
TELECOMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAX: 206587040606
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
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TOPOLOGY: linear
MOLECULE TYPE: CDNA
HYPOTHETICAL: NO
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⁷⁸¹ CTCTGA 786 | | | | | | | | 826 CTCTGA 831

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Search completed: May 30, 2002, 02:51:45 Job time: 11609 sec

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PAT 10-AUG-1998
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AF344841 Cercocebu
AF344859 Macaca mu
X96710 H.sapiens m
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215017 H.sapiens m
X67878 H.sapiens m
L07414 Human CD40-
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248469 B.taurus mR
AF344853 Macaca ne
AF079105 Felis cat
                       AR078316 Sequence
AR085419 Sequence
AR103375 Sequence
AR106246 Sequence
AR169232 Sequence
AR171647 Sequence
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AF344844 Callithri
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Srinivasam.S. and Spriggs,M.K.
Methods of preparing soluble, oligomeric proteins
Patent: US 5716805-A 3 10-FEB-1998;
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Pred. No. 1.3e-188;
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Sequence 3 from patent US 5716805.
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187864.1 GI:3407804
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TITLE
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                                                                             May 30, 2002, 02:49:07; Search time 6499.83 Seconds
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Compugen Ltd
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Pred. No. 1.3e-188;
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Patent: US 5817516-A 3 06-0C5
Location/Qualifiers
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Kehry, M. and Castle, B.
Methods for proliferating and differentiating
density membrane CD40 ligand
           20;
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Sequence 3 from patent US 5817516.
AR044779
             Mismatches
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Armitage,R.J., Fanslow,W.C. and Spriggs,W.K.
Monoclonal antibodies to CD40 ligand, pharmaceutical
Monoprising the same and hybridomas producing the same
Patent: US 5961974-A 11 05-OCT-1999;
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Armitage, R.J., Fanslow, W.C., Spriggs, M.K., Srinivasan, S., Gibson, M.G., Morris, A.E. and McGrew, J.T.
Recombinant soluble CD40 ligand polypeptide and pharmaceutical composition containing the same patent: US 5962406-A 11 05-0CT-1999;
Location/Qualifiers
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Pred. No 1.3e-188;
0; Mismatches 20;
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Sequence 11 from patent US 5962406.
AR078316 AR078316.1 GI:10005062
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Best Local Similarity 97.5%;
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                  CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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Armitage, R.J., Fanslow, W.C. and Spriggs, M.K.
CD40 ligand polypeptide
Patent: US 6087329-A 11 11-JUL-2000;
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No. 1.3e-188;
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         CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
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Method of augmenting a vaccine response by administering
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GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG
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Spriggs,M.K., Armitage,R.J., Fanslow,W.C. III and Wi
Treatment of individuals exhibiting defective CD40L
Patent: US 6106832-A 7 22-AUG-2000;
Location/Qualifiers
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Sequence 7 from patent US 6106832.
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Kehry,M. and Castle,B.
B cell culture system comprising high density membrane ligand
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Location/Qualifiers
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Sequence 3 from patent US 6297052.
AR171647.1 GI:17910597
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            Unclassified.

1 (bases 1 to 840)
Aruffo,A., Hollenbaugh,D. and Ledbetter,J.
Soluble and its use in B cell stimulation
Patent: US 5540926-A 1 30-JUL-1996;
Location/Qualifiers
                                                                                                                                20;
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Pred. No. 1.3e-188;
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Pred. No. 1.3e-188;
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541	Db 586 CGGGAAGCTTC OY 601 TTCGAGAAAT	646	OY 661 CAATCCATTC 	721	781	Db 826 CTCTGA 831		ACCESSION AX090039 VERSION AX090039 VERSION AX090039	SM	Z E	REFERENCE 1 (DASSE AUTHORS Ahuja,S.) TITLE C440 agoi JOURNAL Patent: BOARD OF	FEATURES Source	BASE COUNT 274 ORIGIN	Query Match Best Local Similar Matches 766; Con	Oy 1. ATGATAGAAA 	Oy 61 ATGAAGATTT 	Qy 121 CTTTTTGCTG 	Oy: 181 GAAGATTTTG 	OY 241 TTACTGAACT 	Oy 301 AACAAAGAGG
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706 CAATCCATTCACTTGGGAGGAGTAITTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 765	721 GTGACTGATCCAAGCGAGCCATGGCACTGGCTTCACGTCTTGGCTTGCTAA 780, 111111111111111111111111111111111111	781 CTCTGA 786	826 CICTGA 831	н 0	N 167828 167828.1 GI:2829950	SM	Unclassified. 1 (bases 1 to 84 Armitage, R.J., Fo Method of prevent		source 1.840 /organism="unknown" count 266 a 185 c 175 g 214 t		Query Match 95.9%; Score 754; DB 6; Length 840; Best Local Similarity 97.5%; Pred. No. 1.3e-188; Matches 766; Conservative 0; Mismatches 20; Indels 0; Gaps 0;	1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGIGGCAACTGGACTTCCAGGAGC 60					301 AACAAAGAGGAGGAGGAAGAAAAAAAAGGTTTGAAATGCAAAAAGGTGATCAGAATCCT 360 	361 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 420	421 GCTGAAAAAGGATACTACACCATGAGCAACTTGGTAACCCTGGAAAATGGGAAACAG 480	481 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT 540 %
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                                              NICTIACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
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rity 97.5%; Pred. No. 1.3e-188;
nservative 0; Mismatches 20;
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/db_xref="taxon:9606"
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1 (bases 1 to 879)
Hollenbaugh,D., Grosmaire,L.S., Kullas,C.D., Chalupny,N.J., Braesch-Andersen,S., Noelle,R.J., Stamenkovic,I., Ledbetter,J.A.
and Aruffo,A.
The human T cell antigen gp39, a member of the TNF gene family, is a ligand for the CD40 receptor: expression of a soluble form of EMBO J 11 (12), 4313-4321 (1992)
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The original submission [1] reported 934bp.
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/protein_id="CAA78737.1"
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PRI, Seattle, WA, 98121
3 (bases 1 to 879)
Hollenbaugh, D.
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91ycoprotein 39.
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GQQSIHLGGVFELQPGASVFVNVTDPSQVSHGTGFTSFGLLKL"
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BETKKENSFERGKGDONPOIAAHVISBASSKTTSVLOMBEKCYYTMSNNLYTLENGKO
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                                                                                                            Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1803)
                                                                                                                                                                                        Submitted (28-UL-1992) M. Spriggs, Immunex Res & Development Corporation, 51 University Street, Seattle WA 98101, USA (Deporation, 51 University Street, Seattle WA 98101, USA (Description, 1803) Spriggs, M. K., Armitage, R. J., Strockbine, L., Clifford, K. N., Macduff, B. M., Sato, T. A., Mallszewski, C. R. and Fanslow, W. C. Recombinant human Co40 ligand stimulates B cell proliferation and immunoglobulin E secretion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAAGATITIGIATICAIGAAAACGATACAGAGAIGCAACACAGGAGAAAGAICCITAICC 240
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Pred. No. 1.3e-188;
0; Mismatches 20; Indels. (
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/db_xref="GI:38412"
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/organism="Homo sapiens"
/db_xref="taxon:9606"
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gb_est1:* gb_est2:* gb_htc:* gb_pts:* em_gss:hum:* em_gss_inv:* em_gss_inv:* em_gss_vrt:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

					SUMMARIES	S	
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-	352.6	44.9	492	10	BF599437)) ((((((((((((((((((BF599437 263218 MA
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Ω 4	50.2	6.4	1027	12	A2166561		AZ166561 SP_0088_B
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œ	47.2	6.0	486	σ	AV714252		AV714252 AV714252
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17	47.2	0.9	874	10	BG715241	•	BG715241 602676120

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2 0		10 BI598293 9 AW408255 9 AW372979 10 BE895474 10 BE895474 10 BE8397964 10 BG539029 10 BG539029 10 BG73905 10 BG730477 10 BG720477	ALIGNMENTS	492 bp. 5s taurus cDNA 5'	Cetartlodactyl? Sos. Se,W.M., Freking. White,J., Cho. Laegreid,W.W., Karamycheva,S.,	truction of a ge truction of a ge 7.626-630 (2001) Animal Research enter, NE 68933marc.usda.gov cing. Bases call ptions.	AGGAAACAGCTATGACCAT GTTTTCCCAGTCACGACG TOW: N COLUMN: 5 LOCATION/QUALIFIERS 1. 492 /organism="Bos taurus" /db_xref="taxon:9913"
7 6.0 423 66.0 553 66.0 553 7.9 458 7.9 562 7.9 5.9 773	5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	5 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8		263218 MARC 3BOV BC 263218 MARC 3BOV BC BF599437 GI:1169 EST. COW. BOS taurus EUKATYOta; Metazoa;	utheri vinae; to 49 '', Gro ray,J. on,M.P	racios and con markers and con me Res. 11 (4 80013 tact: Smith TP A, ARS, US Mea Box 166, Clay 167 (24 406) il: smith@mait@mait@mait@mait@mait@mait@mait@mait	PCR PRIMERS FORWARD: AGGAAACAGCTATGACCAT BACKWARD: GTTTTCCCAGTCACGAC Plate: 33 row: N column: 5 Seq primer: ATTTAGGTGACACTAT Location/Qualifiers 1492 //Organism="Bos taur //db_xref="taxon:991"
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Casas,E., Wray,J.E., White,J., Cho,J., Fahrenkrug,S.C., Benneft,
'G.L., Heaton,M.P., Laegreid,W.W., Rohrer,G.A., Chitko-McKown,C.'
Pertea,G., Holt,I., Karamycheva,S., Liang,F., Quackenbush,J. and.
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Cetartiodactyla, Ruminantia, Pecora,
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USDA, ARS, US Meat Animal Research Center
PO Box 166, Clay Center, NE 68933-0166, USA
                                                                                                                                                 DB 10;
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A1982044 638 bp mRNA linear EST 07-MAY-2001 pat.pk0072.c9.f chicken activated T cell cDNA Gallus gallus cDNA clone pat.pk0072.c9.f 5' similar to CD40 ligand, mRNA sequence. A1982044
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Library made from pooled tissue from lymph node, ovary,
fat, hypothalamus, and pituitary."
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                                                          with phred
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC
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Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
Phasianinae; Gallus.
                                                          Single pass sequencing. Bases called and trimmed with p v0.980904.e. Vector identified by cross_match with the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 398,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      301 AACAAAGAGGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 277.2; DB 9
Pred. No. 1.5e-52;
0; Mismatches 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 638)
Tirunagaru,V.G., Sofer,L., Cui,J. and
An expressed sequence tag database of
                      Fax: 402 762 4390
Email: smith@email.marc.usda.gov
                                                                                                                                                               BACKWARD: GTTTTCCCAGTCACGACG
Plate: 34 row: F column: 23.
Seg primer: ATTTAGGTGACACTATAG
                                                                                                                                                                                                                                                                                        /db_xref="taxon:9913"
/clone_lib="MARC 1BOV"
                                                                                                                                                                                                                                                                           /organism="Bos taurus
                                                                                                                                                                                                                                                                                                                                          /tissue_type="pooled"
                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                 FORWARD: AGGAAACAGCTATGACCAT
                                                                                                                                                                                                                                                                                                                                                                /lab_host="DH10B
                                                                                                     and -minmatch 12 options. PCR PRimers
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87.6%;
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402 762 4390
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Pasadena California
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                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches 142;
                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SP_0088_B1_C02_T7A Strongylocentrotus purpuratus, purple sea urchin, sperm genomic BAC library Strongylocentrotus purpuratus genomic clone Plate=88 Col=3 Row=F, DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cameron,R.A., Mahairas,G., Rast,J.P., Martinez,P., Biondi,T.R.,
Swartzell,S., Wallace,J.C., Poustka,A.J., Livingston,B.T., Wray,
.G.A., Ettensohn,C.A., Lehrach,H.; Britten,R.J, Davidson,E.H. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       498 AGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAATCGGGAAGCTTCGAGTCA 557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  558 AGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCGGTAGATTCGAGAGAATCTTACT 617
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        57 -GCGCCATTCACCCTCTATATTTATTGTACCTCCCCATGGAAGAGGACCGGCTCCTGAT 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             618 CAGAGCTGCAAATACCCACAGTTCCGCCAAACCT---TGCGGGCAACAATCCATTCACTT 674
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAATGTGACTGATCCAAG 734
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              176 GGCCGGTGTCTTCGAGCTGCCGCAGGCCATGTCTTTGTCAATGTGACGGACTCAAC 235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 AGGCTCTACTACTACTACTCACAAGTCAGCTTCTGCACCAAGGCGGCGCGCTTCG---- 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Strongylocentrotus purpuratus
Eukaryota; Metazoa; Echinodermata; Eleutherozoa; Echinozoa;
Echinoidea; Eucchinoidea; Echinocea; Echinoida;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        additional resources
Proc. Natl. Acad. Sci. U. S. A. 97 (17), 9514-9518 (2000)
                                                                                                                                                                                                                                                                                                                                                                                                                           Length 638;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          236 AGCAGTGAACGTCAACCTGGCAACACCTACTTTGGCATGTTCAAGCTGT 285
                                                                                                                                                                                                                                                                                        /cell_type="Con A-activated splenic T cell"
/lab_host="E_coll TOP10 F'"
/note="Vector: pcDNA3"
171 c 151 g 180 t 4 others
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                                                                                                                                                                                                           /db_xref="taxon:9031"
/clone="pat.pk0072.c9.f"
/clone_lib="chicken activated T cell cDNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Contact: Cameron, RA, Davidson, EH, Hood, L
                                University of Delaware
40 Townsend Hall, Newark, DE 19717, USA
Tel: 302 831-1345
Fax: 302-831-3411
Seq primer: T7.
                                                                                                                                                                                                                                                                                                                                                                                                                           Score 69.6; DB 9;
Pred. No. 1.2e-05;
0; Mismatches 114;
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California Institute of Technology
                                                                                                                                                                                             /organism="Gallus gallus"
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                                                                                                                                                       Location/Qualifiers
                    Molecular Endocrinology
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  Contact: Joan Burnside
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ilarity 57.6%;
Conservative (
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Matches 167; C
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Drosophila melanogaster genome survey sequence TET3 end of BAC # BACKI2K22 of RPCI-98 library from Drosophila melanogaster (fruit AL060767
                                                                                                                                                                                                                                                                                                 /organism="Strongylocentrotus purpuratus"
//db.xref="taxon:7668"
//clone="plate="88 Col=3 Row=F"
//clone_lib="Strongylocentrotus purpuratus, purple sea
//clone_lib="Strongylocentrotus purpuratus, purple sea
//clone_lib="Strongylocentrotus purpuratus, purple sea
//clone_lib="Sperm genomic BAC library"
//note="Organ: sperm; Vector: BACe3.6; BAC Clones in E-Coli
DH108:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Direct Submission
Submitted (02-JUN-1999) Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - FRANCE (E-mail : segref@genoscope.cns.fr
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              261 TAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTAAACAAAGAGGAGAGGAGAAGAA 320
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                201 AACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCCTTACTGAACTGTGAGGAGAT 260
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           967 AAGAAAAAAAGGAGAAAAGATCAAAAAATATATATGTTTATAGTGATGGGAAAAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   381 AAGTGAGGCCAGCAGTAAAACAACATCTGTTTACAGTGGGCTGAAAAAGGATAC 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  787 AAGAAAAAAAAAAAAAAAAAAATAAAACAGAACTAGAAACGCAAAAAGAGAC 733
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Muscomorpha; Ephydroidea; Drosophilidae; Drosophila
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 1027;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        6.4%; Score 50.2; DB 12; 48.1%; Pred. No. 0.3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Mismatches 153;
                                                                                                                              Seq.primer: T7
Class: BAC ends
High quality sequence stop: 1027.
. Location/Qualifiers
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                                                               Email: acameron@caltech.edu
Plate: 88 row: F column:
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Tel: (626) 395-8421
Fax: (626) 793-3047
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Submitted (23-JUL-1999) Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - FRANCE (E-mail : segref@genoscope.cns.fr
              NY. The library is named RPCI-98 and was constructed by partial EcoRI digestion of Drosophila DNA provided by the BDGP from the isogenic strain y2; cn bw.sp, the same strain used for the BDGP's, pl and EST libraries. A more detailed description of the library and how to order individual BAC clones, the entire library, or filters for hybridization from the BACPAC Resource Center can be found at http://bacpac.med.buffalo.edu/drosophila_bac.htm.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Drosophila melanogaster (fruit
Park Cancer Institute in Buffalo,
38 and was constructed by partial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 - Web : www.genoscope.cns.fr)
Determination of this BAC-end sequence was carried out as part
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     380 TAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGGGCTGAAAAAAGGATACTACA 439'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             200 AAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCCTTACTGAACTGTGAGGAGA 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATAGAAGGTIGGACAAGAIAGAAGAIGAAAGGAAICIICAIGAAGAIITIGIAIICAIGA 199
                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Inse
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
1 (bases 1 to 1043)
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                                                                                                                                                              /organism="Drosophila melanogaster"
/db_xref="taxon:7227"
/clone_lib="RPCI-98"
/clone="BACR12K22"
                                                                                                                                                                                                                                             538
                                                                                                                                                                                                                                                                                                           DB 12;
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BACN1G11 of DrosBAC library from I
fly, genomic survey sequence.
AL103735
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mRNA linear EST 13-JAN-2001
Homo sapiens cDNA; mRNA sequence.
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                                                     d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prépared from embryos by Alain Bucheton and Genevieve Payan. It has been constructed in the vector.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              181 GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC
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This Drosophila melanogaster
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    collaboration with the European Drosophila Genome Pro
http://www.edgp.ebi.ac.uk . This Drosophila melanoga
library (Dros BAC) was made by Alain Billaud at CEPH
                                                                                                                                                                                                                                                                                                            167 others
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ilarity 35.1%; Pred. No. 0.76;
Conservative 51; Mismatches
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/organism="Drosophila
/plasmid="pBeloBAC11"
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20202663
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Laboratory of Cancer Genetics
Ludwig Institute for Cancer Re
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                                                                                                                                                                                                                           /db_xref="taxon:7227"
/clone_lib="DrosBAC"
/clone="BACN11G11"
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Fax: 86-21-50801922
Email: hanzg@chgc.sh.cn
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AV714252 AV714252.1 GI:10795769
                                                                                                                                                                                                                                                                                                                                                                                                       /note="Organ: Kidney_tumor; Vector: puc18; Site_1: Smal; Site_2: Smal; A mini-library was made by cloning products derived from ORESTES PCR (U.S. Letters Patent application No. 196,716 - Ludwig Institute for Cancer Research) profiles into the pUC 18 vector. Reverse transcription of tissue mRNA and cDNA amplification were performed under low stringency conditions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .;
0
                                                                                                             This sequence was derived from the FAPESP/LICR Human Cancer Genome Project. This entry can be seen in the following URL (http://www.ludwig.org.br/scripts/gethtml2.pl?tl=MR1&t2=MR1-RT0025-171100-003-c06&t3=2000-11-17&t4=1)
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I (bases 1 to 486)
Xu,X., Gu,J., Liu,F., Qu,J., Zhao,M., Li,Y., Huang,Q., Zhou,J., Song,H., Gu,Y., Xang,Y., Gao,G., Xiao,H., Li,N., Qian,B., Gao,X., Cheng,Z., Xu,S., Gu,W., Tu,Y., Jia,J., Fu,G., Ren,S., Zhong,M., Lu,C., Cheng,Z. and Han,Z.
Rua Prof. Antonio Prudente 109, 4 andar, 01509-010, Sao Paulo-SP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Contact: Zeguang Han
Chinese National Human Genome Center at Shanghai
351 Guo Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai
201203, P. R. China
Tel: 86-21-50801919(ex.45)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        238 TCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            298 TTAAACAAAGAGGAGGAGGAAGAAAGAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAT 357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   358 CCTCAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAG 417
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 266 CAGAAAAATGTAGAAGTGAAAACGAAAAAGAACAAACAGAAAACACCTGGAAATGG 321
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6.0%; Score 47.2; DB 10;
50.0%; Pred. No. 1.4;
Live 0; Mismatches 118;
                                                                                                                                                                                                                                                                                                                  /organism="Homo sapiens"
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/clone_lib="RT0025"
                                                                                                                                                                                                    Seq primer: puc 18 forward: High quality sequence start: 8 High quality sequence stop: 396 Location/Oualifiers
                                                                                         Email: asimpson@ludwig.org.br
                                                                                                                                                                                                                                                                                                                                                                                     /dev_stage="Adult"
                                             Tel: +55-11-2704922
Fax: +55-11-2707001
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Matches 118; Conservative
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 578)
Gu,Y., Peng,Y., Song,H., Huang,Q., Yang,Y., Gao,G., Xiao,H., Xu,X., Li,N., Qian,B., Liu,F., Qu,J., Gao,X., Cheng,Z., Xu,Z., Zeng,L., Xu,S., Gu,W., Tu,Y., Jia,J., Fu,G., Ren,S., Zhong,M., Lu,G., Hu,R., Chen,J., Chen,Z., and Han,J., Fu,G., Ren,S., Zhong,M., Lu,G., Hu,R.,
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Chinese National Human Genome Center at Shanghai
Chinese Shoujing Road, Zhangjiang Hi-Tech Park, Pudong, Shanghai
201203, P. R. China
Tel: 86-21-50801919(ex.45)
Fax: 86-21-50801992
                                                                                                                                                                                                      /cell_type="dendritic cells"
/dev_stage="mature"
/lab_host="BM25.8"
/note="Vector: pTriplEx2; Site_1: sfilA; Site_2: sfilB"
/ 7 c 124 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AV724813 HTB Homo sapiens cDNA clone HTBAKC04 5', mRNA sequence. AV724813
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This clone is available at CHGC in Shanghai.
This clone is available at CHGC in Shanghai.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 6.0%; Score 47.2; DB 9; Best Local Similarity 50.0%; Pred. No. 1.4; Matches 118; Conservative 0; Mismatches 118;
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/dev_stage="Adult"
/lab_host="SOLR"
                                                   1. .486 /organism="Homo sapiens" /db_xref="taxon:9606" /clone="DCBADFO7" /clone=lib="DCB"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens'
/db_xref="taxon:9606"
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Unpublished (2000)
                              Location/Qualifiers
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NotI; Site_2: XhoI; cDNA made by oligo-dT priming.
Size-selected on agarose gel. Average insert size -1kb; 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Melton, D., Brown, J., Kenty, G., Permutt, A., Lee, C., Kaestner, K., Lemishka, I., Scearce, M., Brestelli, J., Gradwohl, G., Clifton, S., Hillier, L., Marra, M., Pape, D., Wylle, T., Martin, J., Bittain, A., Schmitt, A., Theising, B., Ritter, E., Ronko, I., Bennett, J., Cardenas, M., Gibbons, M., McCann, R., Cole, R., Tsagareishvill, R., Williams, T., Jackson, Y. and Bowers, Y.

Endocrine Pancreas Consortium
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Dept of Molecular and Cellular Biology, 7 Divinity Ave; Cambridge,
MA 02138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammala; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 583)
Site_1: EcoRI; Site_2:
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                                                                                                                                                                                                                                                                                                 88 CICAAATACGIGGACACCAAITIGCAGAAACAGCGAGAACTICAAAAAGCCAATCAGGAG 147
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Contact: Douglas Melton, Klaus H. Kaestner, & Hiroshi Inoue
                                                                                                                                                                                                                                                                                                                                                                                                                                     418 IGGGCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGG 473
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                                                                                                                Length 578;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /tissue_type="Purified pancreatic islet"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BI713130 583 bp mRNA ie02f03 yl HR85 islet Homo sapiens CDNA 5' 095899 UNKNOWN; mRNA sequence.
    /note="Vector: pBluescript sk(-);
                                                                                                                                                        0; Mismatches 118;
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                                                                                                                  6.0%; Score 47.2; Di
50.0%; Pred. No. 1.4;
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Endocrine Pancreas Consortium
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Other_ESTs: ie02f03.x1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (hinoue@im.wustl.edu)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             617-495-1812
                                              2
86
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                                                                                                                                                              Conservative
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BF791160 586 bp mRNA linear BST 12-JAN-2001
602251255F1 NIH_MGC_81 Homo sapiens cDNA clone IMAGE:4338636 5',
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/clone="IMAGE:438656"
/clone="IMAGE:438656"
/clone=lib="NHH MGC_81"
/lab_host="DH10B (T1 phage-resistant)"
/note="Organ: muscle (skeletal); Vector: pDNR-LIB
/note="Organ: muscle (skeletal); Vector: pDNR-LIB
(clontech): Site_1: Sfil (gccgcctcggcc); Site_2: Sfil
(gccattatggcc): 5; and 3' adaptors were used in cloning
(ggccattatggcc): 5; and 3' adaptors vere used in cloning
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Amplified once. Contact information: Hiroshi Inoue, MD, Metabolism Div. (Alan Permutt Lab), Washington University School of Medicine, Box 8127, 660 South Buclid Ave., St. Louis, MO 6310, E-mail: hinoue@imgate.wustl.edu, Tel: 314-362-1916, Fax: 314-747-2692."
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Mammālia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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National Institutes of Health, Mammallan Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     358 CCTCAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAG 417
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs_r@mail.nih.gov
Tissue Procurent: CLONETECH Laboratories, Inc.
cDNA Library Preparation: CLONETECH Laboratories, Inc.
cDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                298 ITAAACAAAGAGGAGGAAGAAGAAAACAGCITTGAAATGCAAAAAGGTGATCAGAAT
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                                                                                                                                                                                                                                                                                                               Pred. No. 1.4;
0; Mismatches 118;
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/db_xref="taxon:9606"
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                                                                                                                                                                                                                                                                                           Score 47.2;
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plate: LLCM1214 row:
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Best Local Similarity 50.03
Matches 118; Conservative
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/clone_lib="LTI_FL013_FBrn1"
/clone_lib="LTI_FL013_FBrn1"
/clone_lib="LTI_FL013_FBrn1"
/clone_lib="pooled tissue from post conception fetuses (20 week, 24 week and 26 week)"
/lab_host="DH10B"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note="Organ: Fetal brain, Vector: pCMVSFORT 6; 1st strand clone as primed with a NotL-oilgo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              pcMvSPORT 6 vector. Library was constructed by Life
Technologies. Contact: Feng Liang Life Technologies, a
division of Invitrogen 9800 Medical Center Drive Rockville
, Maryland 20850, USA Fax: (1) 301 610 8371 Email:
fliang@lifetech.com URL:
http://fulllength.invitrogen.com"
inserts by PCR. This library was enriched for full-length clones and was constructed by Clontech Laboratories (Palo
                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. I (bases: 1 to 797)
Li, W. B., Gruber, C., Jessee, J. and Polayes, D.
Full:length cDNA libraries and normalization
Unpublished (2001)
                                                                                                                                                                                                                                                                      358 CCTCAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACAACATCTGTGTTACAG 417
                                                                                                                                                                                                                                                                                                                                       TCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297
                                                                                                                                                                                                     298 TTAAACAAAGAGGAGGAGGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAT 357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ALS34423 LTI_FL013_FBrn1 Homo sapiens cDNA clone CSODF004YD24 ALS34423 ALS34423
                                                                                                                                                Gaps
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Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - France
Email: seqref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                     418 TGGGCTGAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGG 473
                                                                                                                                                                                                                                                                                                                                                                                                       ;
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                                                                                                              Length 586;
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                                                                                                             Score 47.2; DB 10;
Pred. No. 1.4;
                                                                                                                                               0; Mismatches 118;
                                              118 t
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /organism="Homo sapiens"
/db_xref="taxon:9606"
                                               6
                                               163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AL534423.1 GI:12797916
                                                                                                              6.0%;
50.0%;
                              o, CA)."
                                                                                                                                                Conservative
                                  Alto,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens
                                                                                                              Query Match
Best Local Similarity
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                                               198
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Matches 118;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE
AUTHORS
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COMMENT
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Length 797

Score 47.2; DB 9; Pred. No. 1.4;

Query Match 6.0%; Best Local Similarity 39.6%;

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/clone_lib="NIH_MGC_96"
//issue_type="hypothalamus"
/lab_host="DH10B"
/note="Organ: brain; Vector: pBluescriptR (modified
pBluescript KS+); Site_l: BamHI; Site_2: SalI-XhoI (gtcgag
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BG715238 845 bp mRNA linear EST 08-MAY-2001 602676117F1 NIH_MGC_96 Homo sapiens cDNA clone IMAGE:4798372 5',
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Contacts: Robert Strausberg, Ph.D.
Email: cgapbs-remail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
CDNA Library Preparation: Michael J. Brownstein (NHGKI), Shiraki
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC clone distribution information can be
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              446
                                                                                                                                                                                                                                                                                                                                                                                                          327 CAGCTTTGAAATGCAAAAAGGTGATCAGAATCCTCAAATTGCGGCACATGTCATAAGTGA 386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          483
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           543
GACAAGATAGAAGATGAAAGGAATCTTCATGAAGATTTTGTATTCATGAAAACGATACAG 210
                                 244 GAAAWAAAAAAAAWTTWAAAAAAAGAAAAATTAAATGAAAAAAAGGAAAAAATAAAAGGA 303
                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleosto
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 845)
NHH-MGC http://mgc.nci.nih.gov/.
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           211 AGATGCAACACAGGAGAAAGATCCTTATCCTTACTGAACTGTGAGGAGATTAAAAGCCAG
                                                                                                                                                                                                 304 AAATRTAAAAAAAAAAAAAAATTWAWTTWAWRRAGGATAAAAATAAAAATGRA
                                                                                                                                                                                                                                                                         387 GGCCAGCAGTAAAACAACATCTGTGTTACAGTGGGCTGAAAAAGGATACTACACCATGAG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            447 CAACAACTTGGTAACCCTGGAAAATGGGAAACAGCTGACCGTTAAAAGACAAGGACTCTA
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Plate: LLAM10686 row: b column: 05
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Location/Qualifiers
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/clone="IMAGE:4798372"
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BG715238
BG715238.1 GI:
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(gtcgag
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603249012F1 NIH_MGC_96 Homo sapiens cDNA clone IMAGE:5300787 5',
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Email: cgapbs remail.nih.gov
Tissue Procurement: Miklos Palkovits, M.D., Ph.D.
CDNA Library Preparation: Michael J. Brownstein (NHGRI), Shiraki
Toshiyuki and Piero Carninci (RIKEN)
CDNA Library Arrayed by: The I.M.A.G.E. Consortium (LLNL)
DNA Sequencing by: Incyte Genomics, Inc.
Clone distribution: MGC.clone distribution information can be
); Oligo-dr primed using primer 5 -THTHTHITHITHING size-selected for average insert size 2.3 kb and normalized to Ror 5. This is a primary library enriched for full-length clones and constructed using the for full-length clones in remarration) Library
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                                                                                            Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRL, National Institutes of Health). Note: this is a NIH_MGC Library. a 151 c 228 g
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National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                           177 TTCATACATTACAGTGGTTGGAATAAAATCGGGATGAATGGGTTCCGGAGGAGCAGGTA 236
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 297 CAGTATGCAGAGGGGAÁGATGAGAGGGGCTGCCCCAGGAAAGAAGAAGAAGATCTGGTCTGCAA 356
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          found through the I.M.A.G.E. Consortium/LLNL at: http://image.llnl.gov
Plate: LLAM11761 row: h column: 04
High quality sequence stop: 740.
Location/Qualifiers
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                                                                                                                                                                                                                                                              DB 10;
                                                                                                                                                                                                                                                                                                      Mismatches 118;
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/lab_host="DH10B"
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/db_xref="taxon:9606"
                                                                                                                                                                                                                                                            Score 47.2;
Pred. No. 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BI601432.1 GI:15494371
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Best Local Similarity 50.0%;
Matches 118; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               mRNA sequence.
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                                                                                                                                                                              BASE COUNT
ORIGIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VERSION
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BF5/3/61

861 bp mRNA linear EST 12-DEC-2000

CD0277072F2 NIH_MGC_62 Homo sapiens cDNA clone IMAGE:4251880 5',
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Sfil (ggccactcggcc); Site_: Sfil (ggccattatggcc);
Double-stranded cDNA was prepared from cell line RNA. 5'
Double-stranded cDNA was prepared from cell line RNA. 5'
and 3' adaptors were used in cloning as follows: 5'
and 3' adaptors were used in cloning as follows: 5'
and 3' adaptors were used in cloning as follows: 5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               a, c, or G and N = A, C, G, or T). Average 1.75 kb (range 0.9-4.0 kb). 15/15 colonies inserts by PCR. This library was enriched for
                                                                                                                                                                                                                  ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Average
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           Cap-trapper method (Carninci, in preparation). Library constructed by M. Brownstein (NIMH/NHGRI, National Institutes of Health). Note: this is a NIH_MGC Library."
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NIH-WGC http://mgc.nci.nih.gov/
National Institutes of Health, Mammalian Gene Collection (MGC)
                                                                                                                                                                                                                                                                                                                                                                                                                                    358 CCTCAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACAACATCTGTGTTAAAG 417
                                                                                                                                                                                                                                                                                                                                                                                                                                                              329 CAGIATGCAGAGGGAAGATGAGAGGGGCTGCCCCAGGAAAGAAGACATCTGGTCTGCAA 388
                                                                                                                                                                                                                                                                                                  209 TICATACATTACAGTGGTTGGAATAAAATTGGGATGAATGGGTTCCGGAGGGCAGAGTA 268
                                                                                                                                                                                                                                                                                                                                               298 TTAAACAAAGAGGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAT 357
                                                                                                                                                                                                                                                          238 TCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297
                                                                                                                                                                                                                  Gaps
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Clone distribution: MGC clone distribution information can
                                                                                                                                                                                                                                                                                                                                                                         189 CAGAAAAATGTTGAAGTGAAAAAGGAAAAAGAACAAGAAAACAGGAAATGG 444
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo
clones and constructed using the
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                                                                                                                                                                       Length 845;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CDNA Library Preparation: CLONFIECH Laboratories, cDNA Library Arrayed by: The I.M.A.G.E: Consortium
                                                                                                                                                                                                                      Indels
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                                                                                                                                                                                 DB 10;
                                                                                                                                                                                                                    Mismatches 118;
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High quality sequence stop: 590.
                                                                                                                                                                              Score 47.2; DF
Pred. No. 1.4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Email: cgapbs-r@mail.nih.gov
Tissue Procurement: ATCC/DCTD/DTP
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Unpublished (1999)
Contact: Robert Strausberg,
       for full-length
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full-length clones and was constructed by Clontech Laboratories (Palo Alto, CA)." BASE COUNT 308 a 169 c 220 g 163 t 1 others ORIGIN	Query Match Best Local Similarity 50.0%; Pred. No. 1.4; Matches 118; Conservative 0: Mismatches 118; Indels 0; Gaps	238 TCCTTACTGAACTGTGAGGAGTTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297	158 TICATACATTACAGIGGIIGGAATAAAATCGGGAIGAATGGGTICCGGAGAGCAGAGTA 217	298 TTAAACAAAGAGGAGAGGAAGAAAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAT 357	218 CTCAAATACGTGGACACCAATTTGCAGAAACAGCGAGAACTTCAAAAAGCCAATCAGGAG 277	358 CCTCAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAG 417	278 CAGTATGCAGAGGGGAAGATGAGGGGCTGCCCCCAGGAAAGAAGACATCTGGTCTGCAA 337	418 TGGGCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGG 473	338 CAGAAAAATGTTGAAGTGAAAAAGGAAAAAGAAAACAAAAACAAAAGAAAAGAAAAGA 393
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CDNA of CD40L, a
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                                                                                                            AAQ57984
AAT58122
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Chimeric - Homo sapiens
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09-DEC-1996;
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CD40 ligand gene u
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CD40 Ligand coding
CD40 Ligand gene.
Human CD40 ligand
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                                                                                                                                                                                                  US-08-982-272-7
786
1 ATGATAGAAACATACAGCCA......TTGGCTTACTCAAACTCTGA 786
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/SIDS1/gcgdata/hold-geneseq/genesegn-embl/NA1981.
                                                                                                                             May .30, 2002, 03:03:31; Search time 680.87 Seconds
               GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                    1736436 seqs, 858457221 residues
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Listing first 45 summaries
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AAV39002
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DB

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661 caatccattcacttgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaat

of,

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The present sequence represents an exemplary CD40 ligand gene, comprising nucleotides encoding the extracellular domains (Domains III and IV) of the human CD40 ligand gene (AAV38998) operatively linked to nucleotides encoding the cytoplasmic domain (Domain I) and transmembrane domain (Domain II) of the murine CD40 ligand gene (AAV38997). The sequence is used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of
                                                                                                                                                                                                                                                                                                                                                                                  molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells. Vectors containing the AML genes can be used in gene therapy for treating neoplasia or autoimmune disorders such as rhemmatoid arthritis. They can also be used for vaccination to produce immunity against a virus cell, bacteria, protein, fungus or neoplasia.
                                                                                                                                                                                                                                                                                                                                                             human cells which comprises introducing a gene encoding an accessory
Vectors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatment neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
                                                                                                                   Page 107; 167pp; English
                                                                                                                      Disclosure;
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Sequence 786 BP; 250 A; 166 C; 170 G; 200 T; 0 other;

to produce neoplasia.

420 GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCTGGAAAATGGGAAACAG 480 gctgaaaaaggatactacaccatgagcaacaacttggtaaccctggaaaatgggaaacag 480. 540 009 009 360 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT 540 CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGAAGGAAAGGAATCTTCAT 180 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300 300 AACAAAGAGGAGAGGAAGAAAAAAAAAGGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 360 ATGAAGAITITITATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120 1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC 60 541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 241 ttactgaactgigaggagattaaaagccagtttgaaggctttgtgaaggatataatgtta CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAAACAACATCTGTGTTACAGTGG ; 0 Length 786; Indels Score 786; DB 19; Pred. No. 4.3e-216; Mismatches 6 Best Local Similarity Matches 786; Conserv Query Match 301 361 421 421 481 541 601 241 301 361 121 61 61 121 181 181 Db qq ŏ g δ Qγ g g δλ 셤 Q g pp ò δ ò a õ 엄 ŏ δ

qq

501 ttcgagagaatcttactcagagctgcaaatacccacagttccgccaaaccttgcgggcaa 660

661 0

g

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The present sequence represents an exemplary CD40 ligand gene, comprising nuclectides encoding the extracellular domains (Domains III and IV) and transmembrane domain. (Domain II) of human CD40 ligand gene (AAV38998) operatively. Inliked to nuclectides encoding the cytoplasmic comming to exemplify the murine CD40. Ligand gene (AAV38997). The sequence is used to exemplify the method of the invention. The specification chescribes a method for altering the immoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Vectors containing the AML genes can be used in gene therapy for treating neoplasia or autoimmune disorders such as rheumatoid arthritis. They can also be used for vaccination to produce immunity against a virus and harteris where the second produce immunity against a virus and the second produce immunity.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CD40 ligand; alteration; immunoreactivity; human cell; accessory molecule ligand; AML; gene therapy; treatment; neoplasia; autoimmune disorder; rheumatoid arthritis; vaccine; chimera; ss.
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                       altering immunoreactivity of cells, particularly for treatment of neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 786;
                                                                                                                                                                                                                                                                                                                                                                                                                              Exemplary CD40 ligand gene used in the course of the invention.
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pred. No. 2.5e-213;
0; Mismatches 6;
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Chimeric - Homo sapiens.
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      ATGAAGATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   accessory molecule ligand; AML; gene therapy; treatment; neoplasia; autoimmune disorder; rheumatoid arthritis; vaccine; chimera; ss.
                                                                                 gaagattttgtattcatgaaaacgatacagagatgcaacacaggagaaagatccttatcc
                                                                                                                                                                                                 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                   CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT
                                                                GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAGATCCTTATCC
                                                                                               TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA
                                                                                                                             AACAAAGAGGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                           CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
                                                                                                                                                                                                                        CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                                                                                                                                                        Exemplary CD40 ligand gene used in the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CD40 ligand; alteration; immunoreactivity; human cell;
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Chimeric - Homo sapiens
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The present sequence represents an exemplary CD40 ligand gene, comprising nucleotides encoding the extracellular domains (Domains III and IV) of human CD40 ligand gene (AAV3899) are operatively linked to nucleotides encoding the murine CD40 ligand transmembrane domain (Domain I) (AAV38997) which is operatively linked to the human CD40 ligand gene cytoplasmic domain (Domain I). The sequence is used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells. Vectors containing the AML genes can be used in gene therapy for treating neoplasia or autoimmune disorders such as Theumatoid arthritis. They can also be used for vaccination to produce immunity against a virus cell, bacteria,
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                                                                                                                                                                                                                                                                                                                                                    Vectors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatment neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
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Pred. No. 1.2e-209;
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96US-0032145.
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09-DEC-1996;
18-JUN-1998
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CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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         TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 660
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A'DNA fragment encoding the extracellular (soluble) region of huma CD40-L was ligated to a synthetic oligonucleotide sequence coding for a leader peptide, a 33 amino acid leucine zipper sequence
                                                                                            CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                                                                                                                       CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                                                                                                                                                                                                             protein;
                                                                                                                                                                                                                                                                                                                                                                           Leucine zipper; trimerisation; trimeric CD40-L; fusion protein;
hetero-oligomer; homo-oligomer; type II transmembrane protein;
soluble CD40-L; tumour necrosis factor family; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prepn. of soluble oligomeric mammalian proteins - using host cells to express a fusion protein comprising a leucine zipper domain and a heterologous mammalian protein
                                                                                                                                                                                                                                                                                                                                                         type II transmembrane protein coding sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       t= human_CD40-L
"nucleotides 148-783 code for the
extracellular region (amino acids
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13-AUG-1993;
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(AAR53968) and the Flag (RTM) linker sequence. Cells expressing the fusion construct are grown to accumulate oligomeric, soluble CD40-L in the supernatant. The leucine zipper sequence spontaneously trimerises in solution and fusion proteins comprising the sequence fused to a heterologous mammalian protein also form
                                                                                                                                                                                                                                                       CAAICCAIICACTIGGGAGGAGTAITIGAATIGCAACCAGGTGCTICGGTGITIGICAAI
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                                                                                                             Length 786;
                                                                                                                                   Indels
                                                                              Sequence 786 BP; 250 A; 168 C; 168 G; 200 T; 0 other;
                                                                                                            Score 754; DB 15; ]
Pred. No. 7e-207;
0; Mismatches 20;
                                                                                                               95.98;
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                                                                                                                            Similarity
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RESULT AAV38997

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25-OCT-1991;
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                                                                                                 accessory molecule ligand; AML; gene therapy; treatment; neoplasia; autoimmune disorder; rheumatoid arthritis; vaccine; ss.
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                                                                                                                                                                                                                                                                                                                        Vectors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatment of neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 754; DB 19; Length 786;
Pred. No. 7e-207;
0; Mismatches 20; Indels
                                                                                        human cell;
                                                                CD40 ligand gene used in the course of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 786 BP; 250 A; 168 C; 168 G; 200 T; 0 other;
                                                                                      CD40 ligand; alteration; immunoreactivity;
                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 104; 167pp; English.
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 BP
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Best Local Similarity 97.5%;
Matches 766; Conservative C
AAV38997 standard; DNA; 786
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                                            (first entry)
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Leucine Zipper; fusion protein production; soluble oligomeric protein; heterologous mammalian type II transmembrane protein; activated T cell; heterologous mammalian type I transmembrane protein; antibody production; CD40-L; B-cell proliferation; CD27-L; lymphocyte antigen; ss.
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                                                                                                                                                                                                           CTGACCGTTAAAAGACAAGGAÇTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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910S-0805723.
920S-0969703.
930S-0107353.
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CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA

CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT

TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA

us-08-982-272-7.rng

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This sequence is the coding sequence for the human CD40 ligand (CD40-L). The encoded protein can be used in a fusion protein produced using the method of the invention. The method is for preparing soluble oligomeric protein by culturing a host cell transfected with a vector for the fusion protein. The soluble oligomeric proteins comprise a leucine zipper fused to the N terminus of the extracellular region of a heterologous mammalian type II transmembrane protein or to the C terminus of the extracellular region of a heterologous mammalian type II transmembrane.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          of AAW41171 or AW41172, optionally with conservative amino acid substitutions, provided that the peptide trimerises in solution. A soluble fusion protein comprising the leucine zipper of AAW41171 linked to the extracellular region of CD40. (a type II transmembrane protein campring the leucine zipper of AAW41171 linked to the extracellular region of CD40. (a type II transmembrane protein that is found on activated T cells and acts as a ligand for the B-cell antigen CD40) stimulates B-cell proliferation and antibody production in a similar manner to membrane-bound CD40-L. A soluble fusion protein comprising the leucine zipper of AAW41172 linked to the extracellular region of CD27. (a type II transmembrane protein that binds to the lymphocyte antigen CD27) inhibits binding of CD27c (a fusion protein comprising the extracellular region of CD27 and a human 1961 Fc region)
                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein, where the leucine zipper is a peptide comprising at least part
                                                              proteins - comprising leucine zipper fused transmembrane protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EBV-transformed B cells expressing CD27-L.
                                                                                                                                                                   Example 1; column 19-20; 21pp; English.
                                                                                      proteins
                                                                                   Soluble oligomeric fusion p
to extracellular region of
WPI; 1998-144799/13.
P-PSDB; AAW41178.
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Sequence 786 BP; 250 A; 168 C; 168 G; 200 T; 0 other;

420 AIGAAGATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120 CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGAAGGAAAGGAATCTTCAT 180 cttttgctgtgtatcttcatagaaggttggacaagatagaagatgaaaggaatcttcat 180 aacaaagaggagaggaaagaaagaaaacagctttgaaatgcaaaaaggtgatcagaatcct 360 Gaps 9 9 atgategaaacatacaaccaaacttctccccgatctgcggccactggccatcagc ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG ; 0 Length 786; 95.9%; Score 754; DB 19; ilarity 97.5%; Pred. No. 7e-207; Conservative 0; Mismatches 20; Similarity Query Match Best Local Simi Matches 766; 421 241 421 61 61 121· 121 181 181 241 301 361 361 q qq δý g δy 셤 g 셤 δ qq ò 원 ò à οy à

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This sequence encodes a human CD40-L polypeptide which binds to CD41 CD40-L is a type II membrane polypeptide which has an extracellular region at its C-terminus, a transmembrane region and an intracellular region at its N-terminus. A soluble form of CD40-L lacks the transmembrane domain. -CD40-L activity is mediated by binding with CD40 an dinduces B cell proliferation and induction of antibody secretion, including IGE. Membrane bound CD40-L acts as a
                                                                                                                                                                                                                        transmembrane; region; intracellular; soluble; activity; B cell; proliferation; induction; antibody; secretion; IgE; agonist; antibody; secretion; IgE; agonist;
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                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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                                                                                                  AAQ41506 standard; DNA; 840 BP.
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91US-0805723
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05-DEC-1991;
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                                                                                                                                                                                                            CD40-L DNA.
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hyperigm syndrome
CD40 ligand gene.
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Spriggs MK,
                                                                                                           21-JAN-1994;
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Matches 766;
                                 Synthetic
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      214 T; 0 other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence represents the CD40 ligand gene. Mutations within this gene were identified by the method of the invention. The method comprises isolating DNA from an individual and selectively amplifing the isolated DNA derived from the CD40 ligand gene. The amplification product is then analysed to determine if there is a mutation present and determining if a protein expressed from the ligand gene will bind CD40. The detection of mutations in the CD40 ligand gene allows subsequent treatment of a syndrome resulting in elevated levels of serum IgM and diminished levels of other Ig isotypes, due to mutation in the CD40 ligand gene. ie. X-linked
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Probe; primer; PCR; amplify; polymerase chain reaction; detection; mutation; CD40 ligand gene; IgM; ss.
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Pred. No. 7.2e-207;
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         TICGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                   CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
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differentiation; proliferation; baculovirus; Spodoptera fru
Sf9; insect cell culture; tumour necrosis factor receptor;
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P-PSDB; AAR85486.
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                                                               R85486) that induces long-term
These proliferating B-cells
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proliferation of B cells in vitro or in vivo, partic. differentiated cells
                                                                                                                                       BP; 263 A; 182 C; 181 G; 214 T; 0 other;
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Pred. No. 7.2e-207;
0; Mismatches 20;
                                   Disclosure; Fig 1; 74pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence represents the cDNA sequence of a novel cytokine ligand for CD40 called CD40L is a type II membrane polypeptide, which is expressed by activated CD4+ T cells, and causes B cell proliferation and induction of antibody secretion. The protein can be used to produce monoclonal antibodies, which in turn bind to CD40-expressing cells. This inhibits binding of soluble CD40 to its ligand CD40L. The monoclonal antibody against CD40L is used to inhibit proliferation of neoplastic cells, and is particularly useful in treating B-cell lymphoma (e.g. where induced after transplants or in other cases of immune deficiency such as AIDS), and also melanoma or carcinoma. Since
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    the monoclonal antibodies inhibit neoplastic cells directly, they may not need to be coupled to a toxin or radioisotope, avoiding toxic effects on .
                                                                                                                                                                                                                                       Cytokine; activated CD4+ T cell; CD40; CD40L; monoclonal antibody; neoplastic cell proliferation; B-cell lymphoma; immune deficiency; AIDS; melanoma; carcinoma; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 ATGAAGATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Treating or preventing neoplastic disease associated with CD40-expressing cells - particularly B-cell lymphoma, by administration of CD40-binding protein, preferably antibody
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 840;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 754; DB·18;
Pred. No. 7.2e-207;
0; Mismatches 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Armitage RJ, Fanslow WC, Longo DL, Murphy WJ;
                                                                                                                                                                                                        cDNA of CD40L, a novel cytokine ligand for CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ) US DEPT HEALTH & HUMAN SERVICES
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                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
46.831
/*tag= a
                                                                                                   AAT93782 standard; cDNA; 840
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larity 97.5%;
Conservative
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                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      soluble CD40-ligand
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nes 766; Conserv
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                802 ctctga 807
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Best Local Si
Matches 766,
                                                                                                                                     AAT93782;
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CD40 ligand; TNF receptor family; activated T cell; membrane glycoprotein; cell proliferation; differentiation;

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us-08-982-272-7.rng

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Sequence 840 BP; 266 A; 185 C; 175, G; 214 T; 0 other;
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                                                                                                                                                                                                    method of the invention. The method has been developed for proliferating method of the invention. The method has been developed for proliferating B cells to increase their number at least 100-fold. The method comprises: (a) providing high density, membrane bound CD40 ligand; and (b) culturing one or more B cells in the presence of this ligand. The culture results in a proliferation in the number of B cells of at least induced to differentiate into antibody producing cells in the presence of one or more cytokines. The method can be used for stimulating B-cell. proliferation in vitro or in vivo, e.g. for treating conditions in which B-cell proliferation and activation is suppressed. Eight rounds of division over six days can be achieved, corresponding to a 256-fold increase in cell numbers, which is a vast increase compared to previous.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA.300
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 22 atgategaaacatacaaaccaaacttctccccgatctgcggccactggactgccatcagc
                                                                                                                                                          - by incubating them
                                                                                                                                                                                                                                                                                                                                                                                                                                      1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC
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                                                                                                                                                                                                                                                                                                                                                                  Sequence 840 BP; 263 A; 182 C; 181 G; 214 T; .0 other;
                                                                                                                                                             Increased proliferation of B cells in culture in the presence of membrane-bound CD40 ligand
                                                                                       (BOEH ) BOEHRINGER INGELHEIM PHARM INC
                                                                                                                                                                                           English.
                                      95US-0431055
                                                          95US-0431055
94US-0234580
                                                                                                                                                                                                                                                                                                                                                                                                     Ouery Match 95.9
Best Local Similarity 97.5
Matches 766; Conservative
                                                                                                                                                                                            Example 1; Fig 1; 37pp;
                                                                                                                                 WPI; 1998-556393/47
                                                                                                              Kehry M;
                                                                                                                                                                                                                                                                                                                                                    culturing methods.
                                                                                                                                           P-PSDB; AAW71751
                                                           28-APR-1995;
                                                                      28-APR-1994;
                                       28-APR-1995;
US5817516-A.
                                                                                                              Castle B,
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This sequence encodes the human CD40 receptor ligand (CD40-L). The invention relates to anti-human CD40-L monoclonal antibodies M90 secreted by hybridoma hCD40-M90 (ATCC HB-12055) and M91 secreted by Hybridoma hCD40L-M91 (ATCC HB-12056). M90 and M91 inhibit CD40-L binding to CD40 and the ability of trimeric CD40-L and anti-Immunoglobulin M to induce proliferation of peripheral blood B cells.
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                                                                                                                                                                                                                                                                                           741
                                                                                                                                                                                                                                                                                                                                             541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                                                                                                             TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                                                    gctgaaaaaggatactacaccatgàgcaacaacttggtaaccctggaaaatgggaaacag
                                    CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                     anti-human CD40-L monoclonal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CD40-L; CD40 receptor ligand; anti-human CD40-L monoclonal binding inhibitor; trimeric CD40-L; anti-Immunoglobulin M; peripheral blood B cell; proliferation inhibitor; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAZ27525 standard; cDNA; 840 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human CD40-L coding sequence.
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92US-0969703
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P-PSDB; AAX39938.
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23-OCT-1992;
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                                             Gaps
                                                                                                                  gaagattttgtattcatgaaaacgatacagagatgcaacacaggagaaaagatccttatcc
                                                                                                                                                                                                                                                                                            GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG
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                                    ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC
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                                                                                                         CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT
                                                                                                                                            GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC
                                                                                                                                                      gp39; CD40 ligand; osteoblast cell death; apoptosis; bone loss;
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Length 840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleotide sequence of human gp39 protein, a CD40 ligand
                  Indels
Score 754; DB 20;
Pred. No. 7.2e-207;
                  0; Mismatches
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95.9%;
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                  Conservative
          Similarity
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Query Match
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Matches
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idiopathic autoimmune thrombocytopenia, asthma, a chronic obstructive lung disease, nephrotic/nephritic syndrome, or cancer. They may also be used to treat or prevent bone loss in a subject undergoing, or scheduled for, an organ or bone marrow transplant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present sequence encodes a 9939 protein. It is a CD40 ligand. CD40 ligands are used for reducing osteoblast cell death or apoptosis, and for treating or preventing bone loss in animals, preferably humans, at risk of, or undergoing, bone loss. The bone loss is associated with osteoporosis, osteonecrosis, inflammatory arthritis, post-menopausal osestrogen loss, esterogen loss due to ovariectomy, total historectomy, lupus nephritis, Takayasu's arteritis, Wegener's granulomatosis, anti-glomerular basement membrane nephritis, myositis, scleroderma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CD40 agonist containing composition, used to reduce bone cell death or apoptosis associated with osteoporosis, osteonecrosis and inflammatory arthritis -
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osteoporosis; osteonecrosis; inflammatory arthritis; esterogen loss; ovaritectomy; historectomy; lupus nephitis; Takayasu's arteritis; Wegener's granulomatosis; nephitis; myositis; soleroderma; thrombocytopenia; asthma; lung disease; cancer; ss.
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ilarity 97.5%; Pred. No. 7.4e-207;
Conservative 0; Mismatches 20;
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CD40 ligand; CD40; T cell; T cell receptor; rearrangement; maturation; cell death inhibition; stress-induced; immunosuppressive; anti-thyroid; anti-inflammatory; anti-diabetic; anti-rheumatic; anti-anaemic; opthalmological; anti-psoriatic; nephrotophic; hepatotropic; virucide; dermatological; cytostatic; ss.
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CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG·420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human_CD40_ligand
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rearrangement on T cells can be used to induce T cell receptor rearrangement and enhance T cell affinity for a particular antigen. CD40 engagement can be bought about by contacting CD40 with a CD40-binding agent; e.g. human CD40 ligand. The CD40-binding agents can also be used agent; e.g. human CD40 ligand. The CD40-binding agents can also be used rearrangement, inhibiting environmental stress-induced cell death, altering the specificity of a T cell towards an antigen, inducing T cell death; towards an antigen of enhancing environmental stress-induced cell death (all claimed). T cell affinity maturation towards a specific antigen can be inhibited, especially for a self-antigen in an autoimmune disease, which includes rheumatoid arthritis, usettis, insulin-dependent
                                                                                                                                                                                                                                           diabetes mellitus; haemolytic anaemias, rheumatic fever, Crohn's disease,
Guillain-Barre syndrome, psoriasis, thyroiditis, Grave's disease,
                                                                                                                                                                                                                                                                     myasthenia gravis, glomerulonephritis, autoimmune hepatitis or systemic
lupus erythematosus. Inducing environmental stress-induced T cell death
is carried out in a cancerous T cell or a self-reactive T cell where the
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGGAAAGGAATCTTCAT
                  T cell receptor gene rearrangement for treating autoimm comprises contacting T cells with a CD40-binding agent
                                                                                                                                                                                                                                                                                                                                                                                        Length 1816;
                                                                                                                                                                                                                                                                                                                    environmental stress is a chemotherapeutic agent (claimed)
                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                               Sequence 1816 BP; 520 A; 461 C; 343 G; 492 T; 0 other;
                                                                                                                                                                                                                                                                                                                                                                                         Score 754; DB 21;
Pred. No. 1e-206;
0; Mismatches 20;
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                                                                                                                                                                                                                                                                                                                                                                                            95.9%;
llarity 97.5%;
Conservative
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diseases comprises contact
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Matches 766;
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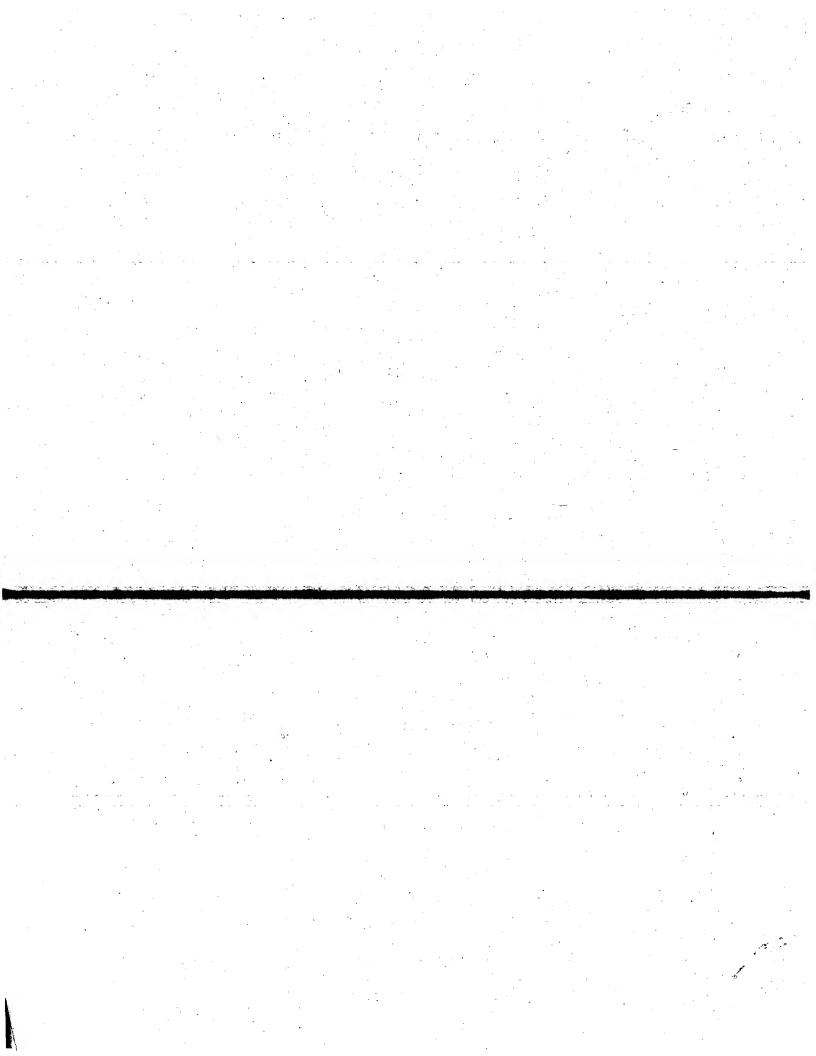
TTCGAGAGAATÇTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGGAA, 660

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The invention relates to isolated polynucleotide (I) and polypeptide (II) sequences. (I) is useful as hybridisation probes, polypeptide (II) sequences. (I) is useful as hybridisation probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags of the for identifying expressed genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving (II). (II) is useful for generating antibodies against it, detecting or quantitating a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful in medical disorders involving aberrant protein expression or biological activity. The polypeptide and polynucleotide sequences have applications in chaptons in cesponsible for genetic disorders or other traits to assess biodiversity
720
                                                                       GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780
                                                                                                                                                and to produce other types of data and products dependent on DNA and amino acid sequences. AAS64197-AAS94564 represent novel human diagnostic coding sequences of the invention.

Note: The sequence data for this patent did not appear in the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; chromosome mapping; gene mapping; gene therapy; forensic; food supplement; medical imaging; diagnostic; genetic disorder; ss.
                                                      CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                                                                                                                                                                                                                                                                                                                                                       novel human diagnostic protein #22375.
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                                                                                                                                                                                                                                                                                                                              AAS86571 standard; cDNA; 1816 BP.
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Copyright (c) 1993 - 2000 Compugen Ltd
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	Sequence	Sequence	Sequence	Sednence	Seguence Seguence	Sequence 1	Sequence	Sequence										e															• 1				٠.,		rth 786;	
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	448-1	993-7 993A-7	926A-	595-3	393-3 631-2: 655-3:	99-30930-1 -447-3848-457	494-1 494A-	926A-	TS					SHOTORY	၂၂၈	144										•													m	
	-430-	545-	-645	-906-	-329- 01-08 01-08	99-30	-09-470-494-1 -09-470-4948-	-09-645-926A-	ALIGNMENT		. ,			V NOT	IC ACCESSORY LIGAND GENES	H			1.				-	, и)	272					. 2								re 786; d. No.	
	ns-09	00-SD	60-SU	60-SN	US-60-329-393-3 PCT-US01-08631-22373 PCT-US01-08655-373	CT-US	60-SN	60-sn	ALI)-)-	272	•	NOTSSEGGAS			CIES		Street				9,00	₹	2.03			2145			13	z.							Score Pred.	
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	32	33	3.5	36 37	39	44.	7 4 4 7 6 4	45			RESULT	; Sequen	GENER GENER	, APP	TITLE OF INVENTION: NOVEL E INVENTION: CONTAIN TITLE OF INVENTION: MOLECULE	TIT	LIL	COF								. coi		, PR.		, AT.		TE		TNFO	SE			90-6	Query	1

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                                                                TITCCITATCACCCAAAIGATIGGATCIGIG 120
                                                                               GGACAAGATAGAAGATGAAAGGAATCTTCAT 180
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                                                                                                             GGACAAGATAGAAGATGAAAGGAATCTTCAT 180
                    CCAGATCCGTGGCAACTGGACTTCCAGCGAGC 60
0; Gaps
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  0; Indels
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FOR IMMUNO-
AND TREATMENT OF
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nucleic acid
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Pred. No. 1.2e-197;
0; Mismatches 6;
                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSeq Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/982,272
                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBRE: 60/132145
FILING DATE: 12/9/96
ATTORNEY/AGENT INFORMATION:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                           MEDIUM TYPE: 3.5" Diskette, MEDIUM TYPE: 1.44 Mb storage
                                                                                                                                                                                                                                                                                          NAME: Guise, Jeffrey W. REGISTRATION NUMBER: 34,613
                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                               TELERA: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS: LENGTH: 786 base pairs TYPE: nucletc acid STRANDEDNES: single
                                                                                                                                                                                                                                                                                                                                                         (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 99.2
Matches 780; Conservative
                                                                                               COMPUTER READABLE FORM:
                                        CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
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                                                                                                                                                                                                                         CLASSIFICATION:
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601 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 660
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                                                                                                                                                                                                                                                                                                                                                       CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                                                                                                                                                                                                                          CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
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APPLICANT: Sharma, Sanjai
APPLICANT: Sharma, Sanjai
APPLICANT: Cantwell, Mark
TITLE OF INVENTION: CONTAINING ACCESSORY
TITLE OF INVENTION: MOLECULE LIGAND GENES AND
TITLE OF INVENTION: MOLECULE LIGAND GENES AND
TITLE OF INVENTION: MODULATION AND TREATMENT OF
TITLE OF INVENTION: MALIGNANCIES
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MODULATION AND TREATMENT OF MALIGNANCIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: 1.44 Mb storage
COMPUTER: 18M Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSeq Version 2.0
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/08/982,272
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NAME: Guise, Jeffrey W.
REGISTRATION NUMBER: 34,613
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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TELECOMMUNICATION INFORMATION:
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ATTORNEY/AGENT INFORMATION:
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California
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OF SEQUENCES:

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CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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                                                                                                                                                            TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
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                                                              Gaps
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                                                                                            541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCGGTAGA
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APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
                                          Length 786;
                                                              Indels
                                      Score 763.6; DB 13;
Pred. No. 3.3e-194;
                                                                0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 3, Application US/08107353 GENERAL INFORMATION:
                                            97.2%;
                                                       Best Local Similarity 98.2
Matches 772; Conservative
 single
STRANDEDNESS: Si
TOPOLOGY: linear
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US-08-107-353-3
                  US-08-982-272-4
                                               Match
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                                                                                                                                                         Version #1.25
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Best Local Similarity 97.5%; Pred. No. 1.2e-191;
Matches 766; Conservative 0; Mismatches 20;
                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: 'IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
                                                                                                                                                                                            US/08/107,353
          ADDRESSEE: Immunex Corporation
STREET: 51 University Street
                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 10
                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION.:
NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,69
                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 786 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: CDNA to mRNA
                                                                                                                                                                                                                                                                                                                  (206)587-0430
                                                                                                                                                                                                                                                                                                                              TELEFAX: (206)233-0644
NFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
                                                                                                                                                                            CURRENT APPLICATION DATA: US
                                                                                                                                                                                                       19930813
                                                                                                    COMPUTER READABLE FORM:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRAIN: CD40-L
                                          Seattle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANTI-SENSE: NO ORIGINAL SOURCE:
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                                                                        COUNTRY:
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RESULT 6
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                                   GCTGAAAAAGGATACTACACCATGAGCAACATGGTAACCCTGGAAAATGGGAAACA 480
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                                                                                                                      CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                          GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG
                                                                       CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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MODULATION AND TREATMENT OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                CONTAINING ACCESSORY
MOLECULE LIGAND GENES AND
                                                                                                                                                                                                                                                                                                                                                                                                                                                       NOVEL EXPRESSION VECTORS
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OPERATING SYZEM: IBM P.C. DOS
SOFTWARE: FastSeq Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MALIGNANCIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/08982272 GENERAL INFORMATION:
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MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: 1.44 Mb storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Guise, Jeffrey W. REGISTRATION NUMBER: 34,613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 23
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/13
FILING DATE: 12/9/96
ATTONEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                Sharma, Sanjai
Cantwell, Mark
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TITLE OF INVENTION:
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California
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781 CTCTGA 786
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STATE:
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                                                                                                                                                                                         Indels
                                                                                                                                                          ; DB 13;
1.2e-191;
                                                                                                                                                                                         Mismatches
                                                                                                                                                          Score 754;
Pred. No. 1
                                                                                                                                                                                      ;
                                                                                                                                                        95.9%;
             INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 786 base pairs
TYPE: nucleic acid
                                                                                                                                                        Query Match 95.9
Best Local Similarity 97.5
Matches 766; Conservative
                                                                         STRANDEDNESS: single
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67-3510
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CAAICCAIICACIIGGGAGGAGIAIIIGAAIIGCAACCAGGIGCIICGGIGIIIGICAAI 720
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601 ITCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
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GENERAL INFORMATION:
APPLICANT: ARMITAGE, MILLIAM
APPLICANT: FANSLOW, WILLIAM
APPLICANT: LONGO, DAN L.
APPLICANT: MIRCHY, WILLIAM
TITLE OF INVENTION: METHOD OF PREVENTING
TITLE OF INVENTION: DISEASE CHARACTERIZED BY NEOPLASTIC CELLS
WINNER OF INVENTION: EXPRESSING CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPANY NATIONAL SYSTEM: Apple Macintosh System 7.1
SOFTWARE: Microsoft Word for Macintosh, Version #5.1a
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STREET: 51 UNIVERSITY STREET
CITY: SEATTLE
STATE: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             December 21,
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MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION DATA:
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826 CTCTGA 831
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           181 GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC 240
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Pred. No. 1.3e-191;
); Mismatches 20; Indels
                                                                                                                                 APPLICANT: SPRIGGS, MELANIE
APPLICANT: WIDMER, MICHAEL
TITLE OF INVENTION: DETECTION AND TREATMENT OF MUTATIONS
TITLE OF INVENTION: IN A CD40 LIGAND GENE
NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                                                                                                                                                                      SYSTEM: Apple Operating System 7.1 MS Word for Apple 5.1, Version a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
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APPLICATION NUMBER: PCT/US94/00786
                 7, Application PC/TUS9400786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/009,258
FILING DATE: 01/22/93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                         E: IMMUNEX CORPORATION 51 UNIVERSITY: STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: PERKINS, PATRICIA ANNE REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 2810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 95.9%;
Best Local Similarity 97.5%;
Matches 766; Conservative
                                                                                                                                                                                                                                                                                                                                                             COMPUTER: Rioppy disk COMPUTER: Apple Macintosh OPERATING SYSTEM: Apple OF
                                                                             DAVISON, BARRY
FANSLOW, WILLIAM
                                                                                                                  RENSHAW, BLAIR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 840 base pairs
TYPE: nucleic acid
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IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2065870606
                                                         APPLICANT: ARMITAGE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             46..831
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: CDNA uvporherICAL: NO
                                                                                                                                                                                                                                                                                                   WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                                       USA
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                                                                                                                                                                                                                                                                                                                                    98101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
LOCATION:
PCT-US94-00786-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ropology:
                                                                                                                                                                                                                                                                                                                   COUNTRY:
                                                                                              APPLICANT:
APPLICANT:
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Query Match

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CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
                                                                                            GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780
                                                                                                                                                                     601 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 660
                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Immunex Corporation
TITLE OF INVENTION: METHOD OF ACTIVATING DENDRITIC CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SYSTEM: Apple Operating System 7.6 Microsoft Word for Macintosh, Version #6.0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION UNBARE:
FILING DATE: 10-JUL-1996
PRIOR APPLICATION DATA:
APPLICATION UNBARE:
APPLICATION UNBARE:
FILING DATE: 12-DEC-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           омыек: PCT/US97/11956
09-JUL-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Power Macintosh 7200/90
                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application PC/TUS9711956 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS: ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION
TELEPHONE: (206)587-0430
TELEPAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Perkins, Patricia A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 34, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      840 base pairs
nucleic acid
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IMMEDIATE SOURCE:
CLONE: CD40-L
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APPLICATION NUMBER: PC
FILING DATE: 09-JUL-19
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PCT-US97-11956-1
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PCT-US97-11956-1
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Pred. No. 1.3e-191;
0; Mismatches 20;
USSN 08/172,664
                 December 23, 1993
                                                 ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34.691
REPERENCE/DOCKET NUMBER: 2818
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 95.9%;
Best Local Similarity 97.5%;
Matches 766; Conservative
                                                                                                                                                                                                                        LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                                                                                                                                                                             STRANDEDNESS: single
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                 FILING DATE: D CLASSIFICATION:
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LOCATION:
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                                                                                                      AIGAAGAITITIAIGIAIITACITIACIGITITICCITAICACCCAAAIGAITGGAICIGG 120
                                                                                                                                                               121. CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
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                                                                                                                                                                                                                                                                                                                286 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 345
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                                            ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC 60
                                                                                                                      GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG
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               0;
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51 UNIVERSITY STREET
 Pred.
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 97.58;
Best Local Similarity 97.5
Matches 766; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                406 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
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                                                                                     SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 754; DB 3;
Pred. No. 1.3e-191;
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                                                                                                                                          US/07/805,723
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0
                                                                    IBM PC compatible
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97.5%;
                                                                                                                                                                                 CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                    Floppy disk
                                                                                                                                                                                                                  NAME: OSTER, JEFFREY B.
REGISTRATION NUMBER: 32
REFERENCE/DOCKET NUMBER:
                                                                                                                         CURRENT APPLICATION DATA: APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                   NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                 COMPUTER READABLE FORM
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                                                                                         SYSTEM:
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766; Conserv
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CLONE: CD40-L
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                                                    MEDIUM TYPE:
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                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                             TELEPHONE:
                                                                      COMPUTER:
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                                                                                         OPERATING
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COUNTRY:
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                                  541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCGGTAGA
                                                  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/969,703A
FILING DATE: 19921023
CLASSIFICATION: 435
ATTONREY/AGENT INPORMATION:
NAME: PERKINS. Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802-B
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1000
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CITY: SEATTLE
STATE: WASHINGTON
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APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 840 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single
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IMMEDIATE SOURCE:
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                                                                                                                                                                                                                                                                                                           CTCTGA 786
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US-07-969-703A-11
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226 GAAGATTTGTATTCATGAAAACGATACAGAGAGATGCAACACAGGAGAAAAAAGATCCTTATCC 285
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                                                  Gaps
                                                                                                                 1 ATGATAGAAACATACAGCCAACCTTCCCCCAGATCCGTGGCAACTGGACTTCCAGCGAGC
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    Length 840,
                                                Indels
  Score 754; DB 3;
Pred. No. 1.3e-191;
                                                0; Mismatches
  Score 754;
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APPLICANT: FNSIGN, WILLIAM
APPLICANT: SPRIGGS, MELANIE
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SOUGHNEES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMINEX CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-07-969-703B-11; Sequence 11, Application US/07969703B; GENERAL INFORMATION:
Query Match 95.9%;
Best Local Similarity 97.5%;
Matches 766; Conservative
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APPLICATION NUMBER: US/08/009,256 FILING DATE: 19930122
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5: IMMUNEX CORPORATION
51 UNIVERSITY STREET
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ELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: PERKINS, PATRICIA A.
REGISTRATION NUMBER: 34,69
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NFORMATION FOR SEQ ID NO:
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MEDIUM TYPE: Floppy
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CLASSIFICATION:
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STATE: WASHING
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STRANDEDNESS
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826 CTCTGA 831
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US-08-009-258-7
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pred. No. 1.3e-191;
0; Mismatches 20; Indels
                                                                                                             OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/969,703B
FILING DATE: 19921023
                                                                               MEDIUM-TYPE: Floopy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                                                                                                        NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
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TELEPHONE: 2065870430
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Best Local Similarity 97.5%;
Matches 766; Conservative
                                                                                                                                                                                                   CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER:
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IMMEDIATE SOURCE:
CLONE: CD40-L
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                       COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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US-07-969-703B-11
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661 CAAICCAITCACIIGGGAGGAGTAITIGAAIIGCAACCAGGIGCIICGGIGIIIGICAAI 720
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                                                  CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: WIDMER, MICHAEL
TITLE OF INVENTION: IN A CD40 LIGAND GENE
TITLE OF INVENTION: IN A CD40 LIGAND GENE
OWNERS PONDENCES: 8
CORRESPONDENCE ADDRESS:
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                                                                                                                       Length 840;
                                                                                                                      Score 754; DB 4; L
Pred. No. 1.3e-191;
0; Mismatches 20;
                                                                                                                    95.98;
                                                                                                                                    Local Similarity 97.5 les 766; Conservative
                                                           46..831
IMMEDIATE SOURCE
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                                          ; NAME/KEY:
; LOCATION:
US-08-009-258-7
                                                                                                                      Query Match
Best Local Si
Matches 766;
                 CLONE:
                            FEATURE:
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TITLE OF INVENTION: METHOD OF PREVENTING OR TREATING
TITLE OF INVENTION: DISEASE CHARACTERIZED BY NEOPLASTIC CELLS
TITLE OF INVENTION: EXPRESSING CD40
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS: ADDRESSE: IMMUNEX CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 840;
                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: Apple Macintosh System 7.1 SOFTWARE: Microsoft Word for Macintosh, Version #5.1a CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/172,664 FILING DATE: December 23, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ouery Match 95.9%; Score 754; DB 5; L
Best Local Similarity 97.5%; Pred. No. 1.3e-191;
Matches 766; Conservative 0; Mismatches 20;
                                                                                                                                                                                       3: IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REPERENCE/DOCKET NUMBER: 2818
TELECOMMUNICATION INFORMATION:
TELEFAX: (206)537-0644
INFORMATION FOR SEQ ID NO: 1:
PANSLOW, WILLIAM
                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
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IMMEDIATE SOURCE:
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                      LONGO, DAN L.
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LOCATION: 46.
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; Sequence 1, Application US/08172664; GENERAL INFORMATION:

US-08-172-664-1

RESULT 13

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GTGACTGATCCAAGCCAAGTGAGCCATGGCATCGCTTCACGTCCTTTGGCTTACTCAAA 780
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                                              CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application US/08215862
GENERAL INFORMATION:
APPLICAMY: Gruss, Hans-J_rgen
TITLE OF INVENTION: Method of Diagnosing or Treating
TITLE OF INVENTION: Disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OPERATING SYSTEM: Apple Macintosh System 7. SOFTWARE: Microsoft Word for Macintosh, Ver CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/215,862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: IMMUNEX CORPORATION STREET: 51 UNIVERSITY STREET CITY: SEATTLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (206)233-0644
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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nucleic acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 424
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US-08-215-862-1
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                                                                                                                     Length 840;
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                                                                                                                                1.3e-191;
                                                                                                                     DB 6;
                                                                                                                                        0; Mismatches
                                                                                                                       Score 754;
Pred. No. 1
                                                                                                                         95.9%;
                                                ORGANISM: Homo sapiens IMMEDIATE SOURCE:
                                                                                                                                          Conservative
                                                                                          46..831
      Linear
                                                                                                                                 Similarity
                        Q.
                                         ORIGINAL SOURCE:
                                                                 CLONE: CD40-L
                       HYPOTHETICAL: N
ANTI-SENSE: NO
                MOLECULE TYPE:
STRANDEDNESS:
TOPOLOGY: li
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Matches 766;
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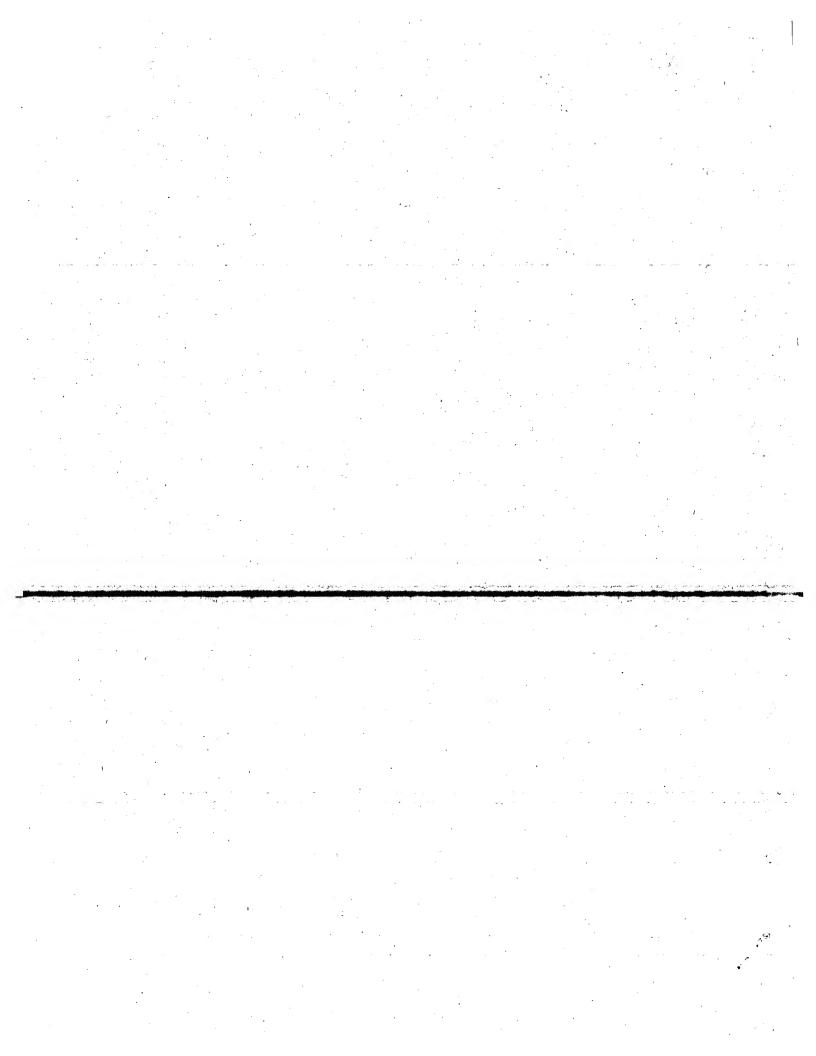
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APPLICANT: CASTLE, BRIAN E
TITLE OF INVENTION: METHODS FOR PROLIFERATING AND
TITLE OF INVENTION: DIFFERENTIATING B CELLS, AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 840;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 754; DB 6; I
Pred. No. 1.3e-191;
0; Mismatches 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1011.1030000/RAM
                                                                                                                                                                                                                                                                                                                                                                                                                   JMBER: US/08/234,580
28-APR-1994
                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                 RESULT 15
US-08-234-580-3
; Sequence 3, Application US/08234580
; GENERAL INFORMATION:
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Best Local Similarity 97.5%;
Matches 766; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MILLMAN, ROBERT A
                                                                                                                                                                                                                                                                                                                                                                                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              840 base pairs
                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
                                                                                                                                                                                         NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nucleic acid
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LOCATION:
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322 AACAAAGAGAGAGAAGAAAAAAAAAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 381
                                                                                                                               GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG 480
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338, ,2, App 4272, A 296495, 208277,

230025, 08276,

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.2937, A .5838, A

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648, App 317028, 317029,

Sequence

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US-10-1027-632-172014
US-10-102-293-172014
US-09-721-544-15838
US-10-027-632-84272
US-10-027-633-296495
US-10-027-633-296495
US-10-027-633-296495
US-10-027-633-296495
US-10-027-633-296495
US-10-027-633-2317029
US-10-027-632-317029
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US-10-027-632-317029
US-10-027-632-45221
US-10-027-632-45221
US-10-027-632-45221
US-10-027-632-45221
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GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
APPLICANT: Chenchik, Alex
TITLE OF INVENTION: Nucleic Acid Arrays
FILE REFERENCE: CLON-006
CURRENT APPLICATION NUMBER: US/09/053,375B
CURRENT FILING DATE: 1998-08-31
                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT FILING DATE: 1998-08-31
NUMBER OF SEQ ID NOS: 1543
SOFTWARE: FastSEQ for Windows Version 4.0
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97.5%;
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US-09-053-375B-226
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Best Local Similarity
Matches 766; Conserv
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LENGTH: 1816
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14649, A
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              GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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US-09-540-210B-14649
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US-09-875-453A-9
PCT-US02-10421-202
US-10-112-699-202
US-09-919-002-664
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-442-384B-457
US-09-053-375B-821
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Listing first 45 summaries
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Maximum DB seq length: 200000000
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120, App 206146,

26090, A 1996, Ap 239, App

206147,

Sequence

US-10-123-155-120 US-10-027-622-206146 US-10-027-632-206147 US-09-539-311D-26090 US-10-105-299-1996

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US-10-027-632-83794 US-10-027-632-83795 US-09-442-384B-628 -09-789-189-1323

628, App 824, App

83795, A

35707, A 62305, A 27253, A

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18 19 20 22 23 25 25 26

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US-10-027-632-62305 US-10-027-632-297617

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Matches 673,
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           caaattgcggcacatgtcataagtgaggccagcagtaaaacaacatctgtgttacagtgg 459
                                                                                  541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 600
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                                         CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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Pred. No. 1.1e-188;
0; Mismatches 20;
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TITLE OF INVENTION: Hemacology/Immunology Array
FILE REFERENCE: CLON-006CIPL5
CURRENT APPLICATION NUMBER: US/09/442,384B
CURRENT APPLICATION NUMBER: US/09/442,384B
PRIOR APPLICATION NUMBER: 099-11-17
PRIOR APPLICATION NUMBER: 099-053,375
PRIOR FILING DATE: 1998.03-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 830
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                 Sequence 457, Application US/09442384B GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95.98;
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Best Local Similarity 97.5
Matches 766; Conservative
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US-09-442-384B-457
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LENGTH: 1816
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CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  GIGACIGAICCAAGCCAAGIGAGCCAIGGCACIGGCIICACGICCIIIGGCIIACICAAA 780
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                                                                                                                                                                                                                                                                                                                               CTGACCGTTAAAAAGACAAGGACTCTATTATATCTATGCCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                               CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAAGTCCCCCGGTAGA
CAAATTGCGGCACATGTCATAAGTGAGGCÇAGCAGTAAAACAACATCTGTGTTACAGTGG
                                                               GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG
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85.6%; Pred. No. 1.2e-146;
1ve 0; Mismatches 110;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Bibilashvilli, Robert TITLE OF INVENTION: Nucleic Acid Arrays FILE REFERENCE: CLON-006 CURRENT APPLICATION NUMBER: US/09/053,375B CURRENT FILING DATE: 1998-08-31 NUMBER OF SEQ ID NOS: 1543 SOFTWARE: FastSEQ for Windows Version 4.0
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; Sequence 821, Application US/09053375B
; GENERAL INFORMATION:
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238 ttacgtctatgcccaagtcaccttctgctccaatcgggcagcttcgagtcaagctccgtt 297
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                                                                                                                                                     ggctagtagtaacccagcgtccgttctgcggtgggcgccaaaagggtactacaccataag
                                                                                                                                                                               CAACAACTTGGTAACCCTGGAAAATGGGAAACAGCTGACCGTTAAAAGACAAGGACTCTA
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Michelotti, Emil F.
Velligan, Mark D.
Latour, Derek R.
Thomas, Rita L.
Kongpachith, Ana
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80.3%;
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Lim, Moon Young
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Starr, Douglas B.
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
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LENGTH: 2395
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APPLICANT:
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APPLICANT:
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APPLICANT:
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                                                                                              GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG
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                           GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAGATCCTTATCC
                                                                                                                                                                                           CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTAACAGTGG
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Pred. No. 2.2e-85;
0; Mismatches 84;
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APPLICANT: Pickett, Gavin G.
APPLICANT: Neft, Robin Eileen
APPLICANT: Dun, II, Robert Thomas
TITLE OF INVENTION: CANINE TOXICITY GENES
FILE REFERENCE: 400742000200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 386
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 131
LENGTH: 508
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Best Local Similarity 83.0%;
Matches 424; Conservative
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US-09-911-904-131
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US-09-911-904-131
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TITLE OF INVENTION: PROMOTERS FOR REGULATED GENE EXPRESSION
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Pred. No. 2.6e-23;
0; Mismatches 36;
                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 54600-8135.US00
CURRENT APPLICATION NUMBER: US/09/875,453A
CURRENT FILING DATE: 2002-05-06
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SOFTWARE: FastSEQ for Windows Version 4.0
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Sequence 9, Application US/09875453A
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Best Local Similarity 80.3
Matches 147; Conservative
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298 TTAAACAAAGAGGAGACGAAGAAAAAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAAT 357
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                                                                                                                                                                                                                                                                                                            298 ITAAACAAAGAGGAGGAAGAAAAAAAAAGAAAAAGGTITGAAATGCAAAAAAGGTGATCAGAAT 357
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                                                                                                                                                                                                                                                                   165 ttcatacattacagtggttggaataaaaattgggatgaatgggttccggagagcagagta 224
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                                                                                                                                                                                                                             TCCTTACTGAACTGTGAGGAGTTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATG 297
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Pred. No. 0.064;
0; Mismatches 120; Indels
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IITLE OF INVENTION: NOVEL CONTIGS OBTAINED FROM VARIOUS CDNA
IITLE OF INVENTION: LIBRARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT FILING DATE: 2001-07-30
CURRENT FILING. DATE: 2001-07-30
PRIOR APPLICATION NUMBER: APPLICATION NUMBER: 09/359,922
PRIOR FILING DATE: FILING DATE: 1999-07-22
DPIOR APPLICATION NUMBER: APPLICATION NUMBER: US 09/034,341
                                                                                                                                                       .DB 6;
                                                                                                                                                  Score 45.6; DB pred. No. 0.017 0; Mismatches
                                                                                                                                                                          0.017
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SOFTWARE: FastSEQ for Windows Version 3.0
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CURRENT APPLICATION NUMBER: US/09/919,002
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GENERAL INFORMATION:
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Best Local Similarity 49.6%;
Matches 117; Conservative
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Best Local Similarity 49.2%
Matches 116; Conservative
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APPLICANT: Liu, Jin
SEQ ID NOS: 2959
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US-09-919-002-664
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    NUMBER OF SEO
SEQ ID NO 202
LENGTH: 569
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          ATGATAGAAACATACAGCGAACCTTCCCCCAGAGCGGGGAACTGGACTTCCAGCGAGC 60
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF CANCER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY TITLE OF INVENTION: AND DIACNOSIS OF CANCER FILE REFERENCE: 210121.565PC
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0; Mismatches 119; Indels
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Pred. No. 0.017;
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CURRENT FILING DATE: 2002-03-20
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CURRENT FILING DATE: 2002-03-28
                                                                                                                                                                                                                                                                                                                                                                                                   Application PC/TUS0210421
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APPLICANT: Wang, Siging
APPLICANT: Bangur, Chaitanya S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bangur, Chaltanya S.
Gaiger, Alexander
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ilarity 49.6%;
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ORGANISM: Homo sapiens
pcr-uS02-10421-202
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Best Local Similarity
Matches 117; Conserv
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US-10-112-699-202
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PCT-US02-10421-202
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                                                                                                    PLICANT: Mullahy, Sara J.
PLICANT: Naughton, Rebesca E.
TLE OF INVERTION: POLYNUCLEOTIDES OF URINARY TRACT TISSUE
LE REFERENCE: PD-1037 CIP
                                                                                                                                                          RRENT APPLICATION NUMBER: US/09/540,210B
RRENT FILING DATE: 2002-04-03
            7000, Application US/09540210B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LING DATE: May 23, 1996
PLICATION NUMBER: 08/881,589
LING DATE: June 24, 1997
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ICATION NUMBER: 60/018,217
                                                                                                                                                                                       OR APPLICATION NUMBER: 08/972, 899
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      08/903,802
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NUMBER: 08/903,471
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ILING DATE: July 31,
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263 AAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTAAACAAAGAGAGGAGGAGAAGAAAG 322
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               121 gggctgccccaggaaagaagacatctggtctgcaacagaaaaatgttgaagtgaaaacga 180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 aaaattgggatgaatgggttccgnagagcagagtactcaaatacgtggacaccaatttgc 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 234;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 39.4; DB 5; Length 2 Pred. No. 0.55; 0; Mismatches 108; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME/KEY: misc_feature
OTHER INFORMATION: Incyte ID No: hu00141818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      443 TGAGCAACACTTGGTAACCCTGGAAAATGG 473
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; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  PLICATION NUMBER: 09/107,592
                                                                                                                                                                TLING DATE: March 12, 1998
PPLICATION NUMBER: 60/040,199
                                                                                                                                                                                                      March 14, 1997
NUMBER: 09/050,817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TLING DATE: July 1, 1997
PPLICATION NUMBER: 09/094,079
                                                                                          60/040,431
APPLICATION NUMBER: 60/039,32
                                                                                                                                                                                                                                                                                                                                                   ILING DATE: May 8, 1998
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APPLICANT: Delegeane, Angelo
APPLICANT: Stuart, Susan G.
                                                                                                                                                                                                                                                        March 30,
                                                                                                                                                             March 12
                          FILING DATE: February
                                                                                                                                                                                                                                                                                                                                                                                                                                         June 30,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 48.8
Matches 103; Conservative
                                                                                          APPLICATION NUMBER: 6
                                                                                                                                                                                                                                                                                                                                                                      LICATION NUMBER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Homo sapiens
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SOFTWARE: PERL PI
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LENGTH: 234
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LICANT: Naughton, Rebecca E.
LE OF INVENTION: POLYNUCLEOTIDES OF URINARY TRACT TISSUE
                                                CURRENT APPLICATION NUMBER: US/09/540,210B CURRENT FILING DATE: 2002-04-03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 08/826,847
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CATION NUMBER: 08/881,589
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DATE: August 22, 1996
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DATE: October 29,
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                                                                                          APPLICATION NUMBER
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238 ICCTIACIGAACIGIGAGGAGAITAAAAGCCAGTIIGAAGGCITIGIGAAGGAIATAAIG 297
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   152 cagtatgcagaggggaagatgagagggctgccccaggaaagaagacatctggtctgca 210
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 38.2; DB 5;
Pred. No. 1.1;
0; Mismatches 88;
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; OTHER INFORMATION: Incyte ID No: hu00678829
US-09-540-210B-14649
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Application US/10123155
          APPLICATION NUMBER: 09/050,817
FILING DATE: March 30, 1998
APPLICATION NUMBER: 60/043,792
FILING DATE: APFIL 11, 1997
                                                                                                                                                                                                               09/094,079
                                                                                 APPLICATION NUMBER: 09/074,999
                                                                                                                 60/048,431
March 14, 1997
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Best Local Similarity 50.8%;
Matches 91; Conservative
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                                                                                                                                                CATION NUMBER:
                                                                                                                                                                                   ICATION NUMBER:
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                                                                                                   NG DATE: May 8,
ICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                 SOFTWARE: PERL Program
SEQ ID NO 14649
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US-10-123-155-120
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PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR PILING DATE: 2000-03-29
PRIOR PELICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-08-09
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Best Local Similarity
Matches 89; Conserv
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LENGTH: 663
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  351 TCAGAATCCTCAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGT 410
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         471 TGGGAAACAGCTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTT 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   591 CCCCGGTAGATTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAAACC 650
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       651 TTGCGGGCAACAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGT 710
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           134 WLRRHAIPDDKLTYGTIAAIGEQNEERLRRLLARPGGGPGGAAQRKVRAFFRSCLDMREI 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GAATCTTCATGAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAG 230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              254 RNSSRYVIRIDQDGLTLPERTLYLAQDEDSEKILAAYRVFMERVLSLLGADAVEQKAQEI 313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            374 VVLLATDYMOQVSQLIRSTPHRVLHNYLVWRVVVVLSEHLSPPFREALHELAQEMEGSDK 433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                554 IRQEVDKSTWLLPPQALNAYYLPNKNOMVFPAGILQPTLYDPDFPQSLNYGGIGTLIGHE 613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATCCTTATCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LQVEQOLANITVSEYDDLKRDVSSMYNKVTLGQLQKITPHLRWKWLLDQIFQEDFSEEEE
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GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 1080827.129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ERLGPRPMLEVIEDCGGWDLGGAEERPGVAARWDLNRLLYKAQGVYSAAALFSLTVSLDD
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                                                                                                                                                                                                                                                                                                                                                                                                                Length 775;
                                                                                                             Prior Application removed - See Palm or File Wrapper NUMBER OF SEQ ID NOS: 550
SEQ ID NO 120
LENGTH: 775
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 45; Conservative 192; Mismatches 404;
                                                                                                                                                                                                                                                                                                                                                                                                                    DB 6;
   TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C30
                                                                                                                                                                                                                                                                                                                                                                                                                    Score 38.2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/027,632 CURRENT FILING DATE: 2002-04-30 PRIOR APPLICATION NUMBER: US 60/218,006
                                                            CURRENT APPLICATION NUMBER: US/10/123,155 CURRENT FILING DATE: 2002-04-15
                                                                                         2002-04-15
                                                                                                                                                                                                                                                                                ; ORGANISM: Homo Sapien US-10-123-155-120
                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-10-027-632-206146
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Best Local S
Matches 45
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176 TTCATGAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAGATCCT 235
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 10827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                296 TGTTAAACAAAGGAGGAGAGGAAGAAAAAACAGCTTTGAAATGCAAAAAGGTG 349
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PRIOR PELING DATE: 2000-07-12
PRIOR PELING DATE: 2000-07-12
PRIOR PELING DATE: 2000-04-20
PRIOR FILING DATE: 2000-04-30
PRIOR FILING DATE: 2000-04-30
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-02-34
PRIOR FILING DATE: 2000-02-34
PRIOR PELING DATE: 1999-11-23
PRIOR PELING DATE: 1999-11-23
PRIOR PELING DATE: 1999-09:28
PRIOR PELING DATE: 1999-09:28
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR PILING DATE: 1999-09:28
NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 206146
LENGTH: 663
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SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                         Pred. No.
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NUMBER
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                                                                                                                                            236 TATCCTTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAA 295
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                                                             176 IICAIGAAGAITITGIAITCAIGAAAACGAIACAGAGAIGCAACACAGGAGAAAAAGAICCI 235.
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                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Naughton, Rebecca E. TITLE OF INVENTION: POLYNUCLEOTIDES OF CARDIOVASCULAR SYSTEM TISSUE FILE BEFERENCE: PD-1022 CIP CURRENT APPLICATION NUMBER: US/09/539,331D CURRENT FILLING DATE: 2000-03-30
                                                                                                                                                                                                                              296 TGTTAAACAAAGAGGAGACGAAGAAAGAAAACAGCTTTGAAATGCAAAAAGGTG 349
                                                                                                                                                                                                                                                                     293 gagatatcagggagctgaaaaatacaatgaataatgtgaaaatgcaatagagag 346
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49.7%; Pred. No. 3.2;
Live 0; Mismatches 90; Indels ©
                      Indels
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                        85;
                          Mismatches
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OTHER INFORMATION: Incyte ID No: hu00341082
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: unsure
LCCATION: 262, 268, 288, 330, 338, 353, 371.
COTHER INFORMATION: a, t, c, g, or other
09-539-331D-26090
    Pred. No. 1.7;
                                                                                                                                                                                                                                                                                                                                                                        Sequence 26090, Application US/09539331D GENERAL INFORMATION:
APPLICANT: Sellhamer, Jeffrey J. APPLICANT: Delegeane, Angelo M. APPLICANT: Stuart, Susan G.
      51.1%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 4.73
Best Local Similarity 49.73
Matches 89; Conservative
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                            Conservative
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NUMBER OF SEQ ID NOS: 4096
SOFTWARE: PERL Program
SEQ ID NO 26090
LENGTH: 438
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ORGANISM: Homo sapiens
    Best Local Similarity
Matches 89; Conserv
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-09-539-331D-26090
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TITLE OF INVENTION: Human Secreted Proteins File Reference: PS950 CURRENT APPLICATION NUMBER: US/10/105,299 CURRENT FILING DATE: 2002-03-26

Sequence 1996, Application US/10105299 GENERAL INFORMATION:

JS-10-105-299-1996/c

APPLICANT: Rosen, et. al

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1174 TAACTGAAGTGTGAAATAACAAAAGAATGCATATAGTCCGTTATTTAAAAATCCCTGTTA 1115
                                                                                                                                                                                                                                                                                                                                                                                                                                                             1114 CACAGAAAAAAAAAAAAAAAAACTTGTTTGAATTTTTAATAGGTCCTAAATCTGGT 1055
                                                                                                                                                                                                                                                                                                                                                                                                                      301 AACAAAGAGGAGGAGGAAGAAAAAAAAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 360
                                                                                                                                                                                                                                                                                                                                      241 ITACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 300
                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                         Length 1751;
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See File Wrapper or Palm
                                                                                                                                                                                                                                                               DB 6;
                                                                                                                                                                                                                                                           Score 36.4; DB Pred. No. 6.1; 0; Mismatches
                                                                                                                                             ; LUCRATION: (1741)..(1742)
; OTHER INFORMATION: n equals a,t,g, or
US-10-105-299-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2002, 05:40:08
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nilarity 53.5%;
Conservative 0
                 SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1996
LENGTH: 1751
  Prior Application removed
                                                                                                       ORGANISM: Homo sapiens
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Best Local Similarity
Matches 76; Conserv
                                                                                      DNA
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(without alignments)
1381.524 Million cell updates/sec
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786
1 ATGATAGAAACATACAGCCA......TTGGCTTACTCAAACTCTGA 786
                                                                                                                     May 30, 2002, 02:51:45; Search time 139.75 Seconds
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                            383533 seqs, 122816752 residues
                                                                             OM nucleic - nucleic search, using sw model
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Gapop 10.0 , Gapext 1.0
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Maximum DB seq length: 2000000000
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Perfect score:
Sequence:
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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

1: /cgn2_6/ptodata/2/ina/5A_COMB.seq:*
2: /cgn2_6/ptodata/2/ina/5B_COMB.seq:*
3: /cgn2_6/ptodata/2/ina/6A_COMB.seq:*
5: /cgn2_6/ptodata/2/ina/eB_COMB.seq:*
5: /cgn2_6/ptodata/2/ina/eB_COMB.seq:*
6: /cgn2_6/ptodata/2/ina/eB_COMB.seq:* Issued_Patents_NA:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	Description	Sequence 3, Appl1	Sequence 3, Appl1	-1	7,	Sequence 1, Appli	3,	H	11,	Sequence 11, Appl	11,	1, 1	11	7,7	11,	Sequence 11, Appl	3, 1	15		Sequence 15, Appl	Sequence 15, Appl	Sequence 15, Appl	15,	10,	20,	20,	Sequence 20, Appl	Sequence 20, Appl	•
SUMMARIES	QI	US-08-446-922-3	PCT-US93-10034-3	US-07-940-605A-1	US-08-184-422-7	US-08-360-923A-1	US-08-431-055-3	US-08-690-096-1	US-08-249-189-11	US-08-484-624A-11	US-08-477-733B-11	US-08-763-995-1	US-09-088-913A-11	US-08-589-771B-7	US-08-769-819-11	US-08-770-974-11	US-08-858-197-3	US-08-249-189-15	US-08-484-624A-15	US-08-477-733B-15	US-09-088-913A-15	US-08-769-819-15	US-08-770-974-15	US-08-446-922-10	US-08-249-189-20	US-08-484-624A-20	US-08-477-733B-20	US-09-088-913A-20	
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20,	Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli Sequence 1, Appli	Sequence 1, Appli Sequence 1, Appli Sequence 5, Appli Sequence 1, Appli Sequence 1, Appli	2222
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US-08-769-819-20 US-08-770-974-20 US-08-446-922-5	US-U8-249-189-1 US-08-484-624A-1 US-08-477-733B-1 US-09-088-913A-1	US-08-769-819-1 US-08-770-974-1 PCT-US93-10034-5 US-08-431-055-1 US-08-858-197-1	US-08-249-189-22 US-08-477-738-22 US-08-477-738-22 US-09-088-913A-22 US-08-769-819-22 US-08-770-974-22
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81.1 81.1 75.6	75.6 75.6 75.6	75.6 75.6 75.6 75.4	56.7 56.7 56.7 56.7 56.7
637.2	594 594 594	594 594 594 592.4 592.4	44455.6 44455.6 44455.6 4455.6 4455.6
30 29	332 332 34	35 337 338 39	40 41 42 44 44 45

ALIGNMENTS

LITLE OF INVENTION:

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GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780
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                                                                                     Gaps
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                                                                                     Mismatches
                                                             Score 754;
Pred. No. 1
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SEMERAL INFORMATION:
APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
                                                           Ouery Match 95.9%;
Best Local Similarity 97.5%;
Matches 766; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TITLE OF INVENTION:
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PCT-US93-10034-3
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; NAME/KEY:
; LOCATION:
US-08-446-922-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 754; DB 5;
Pred. No. 1.2e-209;
0; Mismatches 20;
                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.
                                                                                                                                                                        APPLICATION NUMBER: PCT/US93/10034 FILING DATE:
                        3: Immunex Corporation
51 University Street
                                                                                                                                                                                                                                              34,693
                                                                                                                             IBM PC compatible
                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 1
TELECOMMUNICATION INFORMATION
TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95.98;
                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,69
                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: CDNA to MRNA HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                            TELEFAX: (206)233-0644
FORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                 LENGTH: 786 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
NUMBER OF SEQUENCES: 8
                                                                                                   COMPUTER READABLE FORM:
                                                                                                                                          SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 766; Conserv
                                                                                                                                                                                                                                                                                                                                                                           linear
                                                                                                                                                                                                          CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE: NO ORIGINAL SOURCE:
                                                   Seattle
                                                                            USA
                                                                                                                MEDIUM TYPE:
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                                                                                        98101
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PCT-US93-10034-3
                         ADDRESSEE:
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                                                                                                                                 COMPUTER:
                                                                            COUNTRY:
                                                  CITY: STATE:
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NAME/KEY:
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                          GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG
                                                                               CTGACCGTTAAAAGACÂAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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Patent No. 5540926
GENERAL INFORMATION:
APPLICANT: ARUFFO, ALEJANDRO
APPLICANT: HOLLENBAUGH, DIANE
APPLICANT: HEDBETTER, JEFFREY A.
TITLE OF INVENTION: SOLUBLE LIGANDS FOR CD40
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/07/940,605A FILING DATE: 04-SEP-1992 CLASSIFICATION: 530
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REFERENCE/DOCKET NUMBER: 5624
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212 790-9090
TELEFAX: 212 869-8864/9741
TELEX: 66141 PENNIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION: NAME: Misrock, S. Leslie
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
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STATE: Nor.
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                                                                                                                        ;
0
                                                                                Length 840;
                                                                                                                        Indels
                                                                              Score 754; DB 1;
Pred. No. 1.3e-209;
0; Mismatches 20;
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; Patent No. 556531
; GENERL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
                                                                              Query Match
Best Local Similarity 97.5%;
Matches 766; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: DAVISON, BARRY
22..807
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GIGACTGAICCAAGCCAAGTGAGCCAIGGCACTGGCTTCACGICCTITGGCTTACTCAAA 780
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                                                                                                                                 406 CAAATTGCGCCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
                                                                                                                                                                             GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG
                                                                                                                                                                                                                                             481 CIGACCGITAAAAGACAAGGACTCIATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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                                                                                                                CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
                                                              METHOD OF PREVENTING OR TREATING DISEASE CHARACTERIZED BY NEOPLASTIC CELLS EXPRESSING CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OPERATING SYSTEM: Apple Macintosh System 7.1 SOFTWARE: Microsoft Word for Macintosh, Version #5.1a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: December 23, 1993 CLASSIFICATION: 424
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51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 1, Application US/08360923A Patent No. 5674492 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         December 21, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: ARMITAGE, RICHARD APPLICANT: FANSLOW, WILLIAM APPLICANT: LONGO, DAN L. APPLICANT: MIRPHY, WILLIAM ITTLE OF INVENTION: DISEASE CHATTLE OF INVENTION: EXPRESSING NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk COMPUTER: Macintosh
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MEDIUM TYPE: Floppy
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: SEATTLE STATE: WASHING
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826 CTCTGA 831
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Pred. No. 1.3e-209;
0; Mismatches 20; Indels
                                                        APPLICANT: WIDMER, MICHAEL
TITLE OF INVENTION: DEFECTION AND TREATMENT OF MUTATIONS
TITLE OF INVENTION: IN A CD40 LIGAND GENE
NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                                       COMPUTER: Apple Macintosh
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARRE: MS WORD for Apple 5.1, Version a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/184,422
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2810-A
                                                                                                                                            : IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                     08/009,258
                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: PERKINS, PATRICIA ANNE
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2810
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         95.98;
                                             SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/00
FILING DATE: 01/22/93
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                             Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 840 base pairs
TYPE: nucleic acid
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ADDRESSEE: IMMUNEX C
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Best Local Similarity
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                                RENSHAW
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; LOCATION:
US-08-184-422-7
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Pred. No. 1.3e-209;
0; Mismatches 20;
NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                          TELEPHONE: (206)587-0430
TELEFAX: (206)533-0644
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                            REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                             95.9%;
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IMMEDIATE SOURCE:
CLONE: CD40-L
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Best Local Similarity 97.5
Matches 766, Conservative
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                                                                                                                                                                                                                                                                                                                    APPLICANT: KEHRY, MERILYN R
APPLICANT: CASTLE, BRIAN E
TITLE OF INVENTION: METHODS FOR PROLIFERATING AND
TITLE OF INVENTION: DIFFERENTIATING B CELLS, AND USES THEREOF
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Pred. No. 1.3e-209;
0; Mismatches 20;
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REGISTRATION NUMBER: 36,217
REFERENCE/DOCKET NUMBER: 1011.1030000/RAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/431,055
FILING DATE: 28-APR-1995
CLASSIFICATION: 435
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APPLICATION NUMBER: US 08/234,580
FILING DATE: 28-ARR-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                           US-08-431-055-3
Sequence 3, Application US/08431055
Patent No. 5817516
GENERAL INFORMATION:
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TELEPHONE: (202)371-2600
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INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
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TYPE: nucleic acid
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CURRENT APPLICATION DATA:
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Best Local Similarity 97.5
Matches 766; Conservative
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ADDRESSEE: STERN, KE
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; LOCATION:
US:08-431-055-3
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CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT 180
           361 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
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APPLICANT: HOLLENBAUGH, DIANE
APPLICANT: LEDBETTER, JEFFREY A.
TITLE OF INVENTION: SOLUBLE LIGANDS.FOR CD40
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IBM PC compatible
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COUNTRY: U.C.
10036-2711
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95.9%; Score 754; DB 2; 1
Best Local Similarity 97.5%; Pred. No. 1.3e-209;
Matches 766; Conservative 0; Mismatches 20;
08/069/069/sn
                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/940,605
FILING DATE: 04-SEP-1992.
ATTORNEY/AGENT INFORMATION:
                                                                                                                   NAME: MAGTOCK, S. LOSIIE
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 5624
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212 790-9090
TELEFAX: 212 869-8864/9741
TELEFAX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                             LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
 APPLICATION NUMBER:
FILING DATE: 31-JUL-
CLASSIFICATION: 536
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                                                                                                                                                                                                                                                                                                                                        linear
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; LOCATION:
US-08-690-096-1
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FEATURE
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                                  661 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
                                                                                                 GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 780
622 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 681
                                                                                                              Word for Apple, version 5.1a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Apple Operating System 7.1
                                                                                                                                                                                                                                                                          Sequence II, Application US/08249189
Patent No. 5961974
GENERAL INPORMATION:
APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US/08/249,189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                               E: IMMUNEX CORPORATION 51 UNIVERSITY STREET
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APPLICATION UNMBER: 07/783,707
FILING DATE: October 25, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/POCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2055870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Of
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TYPE: nucleic acid
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
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802 CTCTGA 807
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TELEFAX: 2
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                                                                                                                                                               Length 840
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                                                                                                                                                           Score 754; DB 2;
Pred. No. 1.3e-209;
0; Mismatches 20;
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                                                                                                                                                           95.98;
                                                                                                                                                                                                   Matches 766; Conservative
                                                                               46..831
                                                                                                                                                               Query Match
Best Local Similarity
IMMEDIATE SOURCE:
CLONE: CD40-L
                                                            CDS
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                                                        ; NAME/KEY:
; LOCATION:
US-08-249-189-11
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46 AIGATCGAAACATACAACATTCTCCCCGATCTGCGGCCACTGGACTGCCATCAGC 105
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US-08-477-733B-11
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APPLICATION NUMBER: US/08/484,624A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 840;
                                                                                                                                      APPLICANT: MCGREW, JEFFERY
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 754; DB 2; I
Pred. No. 1.3e-209;
0; Mismatches 20;
                                                  FANSLOW, WILLIAM
SPRIGGS, MELANIE
SRINIVASAN, SUBHASHINI
GIBSON, MARYLOU
                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,733
FILING DATE: June 07, 1995
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
                                                                                                                                                                                                        : IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patitala A.
RECISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802
TELECOMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEPASX: 2065870606
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APPLICATION NUMBER: 07/783,70
FILING DATE: October 25, 1991
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Best Local Similarity 97.5%;
Matches 766; Conservative (
                                  ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                              Apple Macintosh
                                                                                                                      MORRIS, ARVIA E.
                                                                                                                                                                                                                                                                                                                              Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
companients
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
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                                                                                                                                                                        NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                          WASHINGTON
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                                                                                                                                                                                                            ADDRESSEE:
                                                                                    APPLICANT:
APPLICANT:
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                                                                     APPLICANT:
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661 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 720
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61 ATGAAGATTTTTATGTATTTACTTACTGTTTTCCTTATCACCCAAATGATTGGATCTGTG 120
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                                                                                                                                                                              241 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA
                                                                                                                                                                                                                                                                                                                                                                                   361 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
                                                                                                                                                                                                                                                                                                                                                                                                       481 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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                                                                                                                                                      GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGGAGAAAAGATCCTTATCC
                                    106 ATGAAAATTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA
                                                                                                166 CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATGAAGAAGAAGAAGAAGAACTTCAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 11, Application US/08477733B Patent No. 5981724 GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
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226 GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACAGGAGAAAAAAGATCCTTATCC 285
                                                    CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
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                                    241 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATATAATGTTA
                                                                                                            AACAAAGAGGAGACGAAGAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
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TITLE OF INVENTION: METHOD OF ACTIVATING DENDRITIC CELLS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: Microsoft Word for Macintosh, Version #6.0.1
CURRENT APPLICATION DATA:
APPLICATION UNMER: US/08/763,995
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COMPUTER: Power Macintosh 7200/90
OPERATING SYSTEM: Apple Operating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
CITY: SEATTLE
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NAME: Perkins, Patricia
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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FILING DATE: 10 JUL
CLASSIFICATION:
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WASHINGTON
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                                                                                     MEDIUM TYPE: Floppy disk COMPUTER: Apple Macintosh OPERATING SYSTEM: Apple Operating System 7.5.5 SOFTWARE: MS Word for Apple Power Macintosh, version 6.0.1 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/477,733B FILING DATE: June 07, 1995
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                                                                                                                                                                                                               CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/249,189
FILING DATE: May 24, 1994
CLASSIFICATION: 435
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                                                                                                                                                                                                                                                                                                                                         07/969,703
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                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: 07/969,70
FILING DATE: October 23, 1992
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NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,69;
REFERENCE/DOCKET NUMBER: 25
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS: LENGTH: 840 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 95.9%;
Best Local Similarity 97.5%;
Matches 766; Conservative
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APPLICATION DATA:
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                                                                     COMPUTER READABLE FORM:
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CLONE: CD40-L
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                                    USA
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                                                    98101
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US-08-477-733B-11
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                                    COUNTRY:
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721 GIGACIGATCCAAGCCAAGIGAGCCAIGGCACIGGCTTCACGICCTTIGGCTTACTCAAA 780
                                                                 SYSTEM: Apple Operating System 7.5.5
MS Word for Apple Power Macintosh, version 6.0.1
                                                                                                                                                                                                                                                                                                                                                                                            NOVEL CYTOKINE THAT BINDS CD40
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US-09-088-913A-11
. Sequence 11, Application US/09088913A
; Patent No. 6087329
                                                                                                                                                                                                                                                                                                                               SRINIVASAN, SUBHASHINI
GIBSON, MARYLOU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE: June 07, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
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51 UNIVERSITY STREET
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APPLICATION NUMBER: 08/484,624
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 28
ELECOMMUNICATION INFORMATION
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                 ARMITAGE, RICHARD
FANSLOW, WILLIAM
SPRIGGS, MELANIE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                ARVIA E.
                                                                                                                                                                                                                                                                                                                                                                                   MCGREW, JEFFERY
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APPLICATION NUMBER:
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MEDIUM TYPE: Floppy of
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                                                                                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION:
NUMBER OF SEQUENCES:
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WASHINGTON
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826 CTCTGA 831
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                          0;
                                                                                                                                                                                                                                                                                                                                                                                                           Length 840;
                                                                                                                                                                                                                                                                                                                                                                                                           Score 754; DB 3;
Pred. No. 1.3e-209;
0; Mismatches 20;
                                  REFERENCE/DOCKET NUMBER: 284
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEFAX: (206)233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                95.9%;
97.5%;
                                                                                                                                                                                                                                                   ORIGINAL SOURCE:
ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                    NFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                        LENGTH: 840 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                     STRANDEDNESS: single
                      REGISTRATION NUMBER:
                                                                                                                                                      nucleic acid
                                                                                                                                                                                                                                                                                                                            ; NAME/KEY: CDS
; LOCATION: 46..831
US-08-763-995-1
                                                                                                                                                                                                    MOLECULE TYPE: CDNA HYPOTHETICAL: NO
                                                                                                                                                                                   linear
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Best Local Similarity
Matches 766; Conserv
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                                                                                                                                                                                        TOPOLOGY:
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                                                                                                                                                                                                                                                                                                                   FEATURE
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                                                                                                                                        Length 840;
                                                                                                                                     Score 754; DB 3; Pred. No. 1.3e-209; O; Mismatches 20;
                                                                                                                                     Query Match 95.9%;
Best Local Similarity · 97.5%;
Matches 766; Conservative
              IMMEDIATE SOURCE
                              CLONE: CD40-L
                                                          ; NAME/KEY: CDS
; LOCATION: 46..
US-09-088-913A-11
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ORGANISM:
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                                                                                                                             TITLE OF INVENTION: TREATMENT OF INDIVIDUALS EXHIBITING TITLE OF INVENTION: DEFECTIVE CD40L (as amended) NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               95.9%; Score 754; DB 3; I
97.5%; Pred. No. 1.3e-209;
ive 0; Mismatches 20;
                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Word for Windows 95 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: US/08/589,771B
FILING DATE: January 22, 1996
                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: MS-DOS/Windows
                                                                                                                                                                                                         : IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION TELEPHONE: 2065870430 TELEFAX: 2065870606
                   : ARMITAGE, RICHARD
                                                                   APPLICANT: RENSHAW, BLAIR
APPLICANT: SPRIGGS, MELANIE
APPLICANT: WIDMER, MICHAEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: HENRY, JANIS C.
REGISTRATION NUMBER: 34,3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/05/93
FILING DATE: 01/22/93
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 2065870606
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 840 base pairs
TYPE: nucleic acid
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                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                 WASHINGTON
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                                                                                                                                                                                                                           STATE: WASTE
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MOLECULE TYPE: CF
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INFORMATION
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                                                                                                                                                                                                                                                                                                   98101
                                                                                                                                                                                                         ADDRESSEE:
STREET:
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US-08-589-771B-7
                                                                                                                                                                                                                                                                                 COUNTRY:
                                 APPLICANT:
APPLICANT:
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RESULT 13 US-08-589-771B-7 ; Sequence 7, Application US/08589771B

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GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG 480
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                                                                                                                                                                   CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAAACAACATCTGTGTTACAGTGG 420
         Apple Operating System 7.1 CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/769,819
FILING DATE: 19-DEC-1996
CLAS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: SRINIVASAN, SUBHASHINI GERSON, MARYLOO TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40 NUMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 11, Application US/08769819, Patent No. 6264951
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     : IMMUNEX CORPORATION
51 UNIVERSITY STREET.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: ARMITAGE, RICHARD APPLICANT: FANSLOW, WILLIAM APPLICANT: SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
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WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    USA
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
                                                                                                                                                                  FILING DATE: December 5, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
                                                                                                                                                                                                                    FILING DATE: October 25, 1991 ATTORNEY/AGENT INFORMATION: NAME: Perkins, Patricia A.
                                                                                                                                                              December 5, 1991
                                                                                                                                                                                                                                                        NAME: PERKINS, PATTICIA A. REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 281 TELECOMMUNICATION INFORMATION:
                                                     May 24, 1994
APPLICATION NUMBER: 08/4E
FILING DATE: 07-JUN-1995
APPLICATION NUMBER: 08/24
                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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IMMEDIATE SOURCE:
CLONE: CD40-L
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                                                       FILING DATE:
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US-08-769-819-11
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CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 420
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                                                                                                                                                                                                                                                                                                                        46 ATGATCGAAACATACAACATCTCCCCGATCTGCGGCCACTGGACTGCCATCAGC 105
                                                                                                  61 AIGAAGAITITIAIGIAITIACITACCIGIITICCTIAICACCCAAAIGAITGGAICIGIG 120
                                                                                                                           165
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                         Gaps
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                                                                                                               121 CTTTTTGCTGTGTATCTTCATAGAAGGTTGGACAAGATAGAAGATGAAAGGAATCTTCAT
                                                                                                                                                                  GAAGATTTTGTATTCATGAAAACGATACAGAGATGCAACACAGAGAAAAAATCCTTATCC
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                          Indels
Score 754; DB 4; I
Pred. No. 1.3e-209;
0; Mismatches 20;
    95.98;
97.58;
    Query Match 95.9
Best Local Similarity 97.5
Matches 766; Conservative
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                                                                                                                                                                                            TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                  CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: FAMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, WILLIAM
APPLICANT: STRIGS, WILLIAM
APPLICANT: STRINIASAN, SUBHASHINI
APPLICANT: GIBSON, WARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintons
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, version 5.1a
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/770,974
FILING DATE: 20-DEC-1996
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,733
FILING DATE: 02-AUG-1995
APPLICATION NUMBER: 08/249,189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 11, Application US/08770974
Patent No. 6290972
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: 07/969,703
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APPLICATION NUMBER:
FILING DATE: December 5, 1991
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: October 25, 1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: October 23, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   May 24, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: May 24, PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER:
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Pred. No. 1.3e-209;
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              TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                                                                                                                                                                                                                                                                                                                                                                95.9%;
llarity 97.5%;
Conservative
                                                                                                                                                                                                         ANTI-SENSE: NO ORIGINAL SOURCE: OROGANISM: Homo sapiens IMMEDIATE SOURCE: CLONE: CD40-L
REFERENCE/DOCKET NUMBER:
                                                                 INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                     : 840 base pairs
nucleic acid
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TOPOLOGY: linear
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Best Local Similarity
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                                                                                                     LENGTH:
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Search completed: May 30, 2002, 02:51:53 Job time: 11617 sec

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Title: Perfect score:

Sequence:

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Scoring table:

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Minimum Maximum

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AX090039 Sequence
215017 H. sapiens m
X67878 H. sapiens m
L07414 Human CD40-
X68550 H. sapiens T
AF344841 Cercocebu
AF344860 Abcaca mu
X96710 H. sapiens m
AF34484 Callithri
AF013985 Rattus no
AR076918 Sequence
AR078308 Sequence
AR0783411 Sequence
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AR10387 Sequence
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AR171646 Sequence
AX208160 Sequence
Z48469 B. Laurus mR
AF079105 Felis cat
                                                  AR085419 Sequence
AR103375 Sequence
AR106246 Sequence
AR169232 Sequence
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(Assess 1 to 786)

Srinivasan, S. and Spriggs, M.K.

Methods of preparing soluble, oligomeric proteins

Patent: US 5716805-A 3 10-FEB-1998;

Location/Qualifiers
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187864
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TITLE
JOURNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     KEYWORDS
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                                                                                        May 30, 2002, 02:49:14 ; Search time 6499.83 Seconds (without alignments) 2520.909 Million cell updates/sec
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Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                               - nucleic search, using sw model
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AF116582 Rattus no

Sequence

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Seguence Sequence Sequence

M.musculus

X65453

PAT 10-AUG-1998

Length 786;

Score 726.8; DB 6; Pred. No. 2.7e-170;

92.8%; 96.2%;

Query Match Best Local Similarity

Description

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	tches	ATCTGCG(ATCTGCG	TCTTATCA TCTTATCA	TAAGGTCO	ATGCAAC ATGCAAC	TGAAGAC	TGAAGGC TGAAATG	TGAAATG CAGTAAA	CAGTAAA	CTTGGTA 	CTATGCC 	CAGCCTC	CCACAGT	rgcaacca 	CTGGCTTC 				bр 7516.	*		differ	
-	Міѕша	ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 	atgaaaattittatgatttacttactgtttttcttatcacccagatgattgggrcagca 	CTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT	GAAGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAGGAGAAGGATCTTTATCC 	TTGCTGAACTGTGAGGAGTGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA	TTACTGAACTGTGAGGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA AACAAAGAAGAGAAAAAAGAAAAGGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 	AACAGCTŲ SAGGCCAG		GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG 	CTGACCGTTAAAAGACAAGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT 	CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 	TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 	CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 	GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 	· ·		*	840] US 581			ing and ligand	
	e 0;	CAAACTI	TTACTTP 	CATAGAA 	AAAAAGG	ATGAGA	SATTAAA? VAAAGAA? 	SAAAGAA SATAAGTO	ATAAGT	CACCATG	AGGACTC AGGACTC	AGCTCCA AGCTCCA	CAGAGCT	AGGAGTA AGGAGTA	AGTGAGC AGTGAGC				patent	:5966244		to 840) Castle, B. proliferating prane CD40 lig	
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PAT 31-AUG-2000
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                                                                                                      Score 726.8; DB 6;
Pred. No. 2.7e-170;
0; Mismatches 27;
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Seguence 11 from patent US 5961974.
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Patent: US 5817516-A 3 06-OCT-1998;
Location/Qualifiers
                                       /organism="unknown"
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                                                      Armitage, R.J., Fanslow, W.C. and Spriggs, M.K. Monoclonal antibodies to CD40 ligand, pharmaceutical comprising the same and hybridomas producing the same Patent: US 5961974-A 11 05-OCT-1999;
                                                                                                                                                                   Length 840,
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Pred. No. 2.7e-170;
0; Mismatches 27;
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185 c 175 g
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Best Local Similarity 96.2
Matches 756; Conservative
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Armitage, R.J., Fanslow, W.C., Spriggs, M.K., S Gibson, M.G., Morris, A.E. and McGrew, J.T.
Recombinant soluble CD40 ligand polypeptide Composition contraining the same
Patent: US.5962406-A 11 05-OCT-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 726.8; DB 6;
Pred. No. 2.7e-170;
0; Mismatches 27;
  DNA
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patent US 5962406.
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PAT 14-FEB-2001
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                                   CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                       CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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Pred. No. 2.7e-170;
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Armitage,R.J., Fanslow,W.C. and Sprist CD40 ligand Polypeptide
Patent: US 6087329-A 11 11-JUL-2000;
Location/Qualifiers
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286 TTACTGAACTGTGAGGAGTTAAAAGCCAGTTGAAGGCTTTGTGAAGGATATAATGTTA
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                                                       CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                                                                                                                                                                                                                                                                                                                (bases 1 to 840)
Armitage, R.J. Fanslow, W.C., Spriggs, M.K., Srihivasan, S., Gibson, M.G., Morris, A.E. and McGrew, J.T.
DNA encoding CD40 ligand, a cytokine that binds CD40
Patent: US 5981724-A 11 09-NOV-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 840;
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Pred. No. 2.7e-170;
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Sequence 11 from patent US 5981724
AR085419 GI:10012188
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CTCTGA 831
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                 TTGCTGAACTGTGAGAGGAATGAGAAGTTTGAAGACCTTGTCAAGGATATAACGTTA
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Pred. No. 2.7e-170;
); Mismatches 27;
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1 (bases 1 to 840)

Armitage, R.J., Fanslow, W.C., Spriggs, M.K Gibson, M.G.
Method of augmenting a vaccine response ligand
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Location/Qualifiers
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96.2%;
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Best Local Similarity 96.23
Matches 756; Conservative
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Spriggs,M.K., Armitage,R.J., Fanslow,W.C. III and Widmer,M.B.
Treatment of individuals exhibiting defective CD40L
Patent: 18 5106632.2A 7 22-AUG-2000;
Location/Qualifiers
                                                                                                                                                                           CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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Pred. No. 2.7e-170;
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Seguence 7 from patent <sup>1</sup>
AR106246
AR106246.1 GI:12820776
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96.2%;
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Kehry, M. and Castle, B.
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                                                                                                   Length 840;
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                            Ledbetter, J.A.
                                                                                                   Score 726.8; DB 6;
Pred. No. 2.7e-170;
0; Mismatches 27;
                   1 (bases 1 to 840)
Aruffo,A., Hollenbaugh,D. and Ledbetter,J.
Soluble and its use in B cell stimulation
Patent: US 5540926-A 1 30-JUL-1996;
Location/Qualifiers
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Spriggs,M.K., Armitege,R.J. and Fanslow,W.C.
Detection of mutations in a CD40 ligand gene
Patent: US 5565321-A 7 15-OCT-1996;
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Pred. No. 2.7e-170;
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Sequence 7 from patent US 5565321.
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OY 538 CGGGAAGCTTCGAGTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCGGTAGA 597 Db 586 CGGGAAGCTTCGAGTCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 645 Db 588 TTCGAGAGATCTTAGTCAGAGCTGCAAATTCCCAGCTTCGCCGGTAGA 657 Qy 11111111111111111111111111111111111	SULT (090C OCUS CESS CESS	SOURCE ORGANISM Homos appens horacoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Eukaryota; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo. REFERENCE 1 (bases 1 to 879) AUTHORS AUTHORS C440 agonist compositions and methods of use conton to 108-MAR-2001; JOURNAL Patent; WO 0116180-A 108-MAR-2001; BOARD OF REGENTS THE UNIVERSITY OF TEXAS SYSTEM (US) FEATURES 1. 1879 Cource //organism="Homo sapiens"	PASE COUNT 274 a 193 c 190 g 222 t ORIGIN ORIGIN QUEFY MATCH 92.8%; Score 726.8; DB 6; Length 879; Best Local Similarity 96.2%; Pred. No. 2.76-170; Matches 756; Conservative 0; Mismatches 27; Indels 3; Gaps 1; Matches 756; Conservative 0; Mismatches 27; Indels 3; Gaps 1; OY 1 ATGATCGAACATACAACTTCTCCCCCAATTGGGGCCACTGGACTGCCCATCAGC 60 ON 22 ARGATCGAACATACAACAACATCCCCGAACTGGGGCCACTGGGCCATCAGC 81	61 82 82 121 142 142	202 , 241 , 241 , 262 301 , 322
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases, 1 to 879)
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Braesch-Andersen,S., Noelle,R.J., Stamenkovic,I., Ledbetter,J.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             and Aruffo, A.

The human T cell antigen gp39, a member of the TNF gene family, a ligand for the CD40 receptor: expression of a soluble form of gp39 with B cell co-stimulatory activity
EMBO J 11 (12), 4313-4321 (1992)
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/organism="Homo sapiens"
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PRI, Seattle, WA, 98121
3 (bases 1 to 879)
Hollenbaugh, D.
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91ycoprotein 39.
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Hollenbaugh, D.L.
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LTVKRQGLYTYYAQVTFCSNREASSQAPFIASLCLKSPGRFFRILLRAANTHSSAKPC
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Pred. No. 2.7e-170;
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    PRI 06-JUN-1997
                                                                                                                                                                                              Spriggs, M. K., Armitage, R. J., Strockbine, L., Clifford, K. N., Macduff, B. M., Sato, T. A., Maliszewski, C. R. and Fanslow, W. C. Recombinant human CD40 ligand stimulates B cell proliferation and immunoglobulin E secretion
J. Exp. Med. 176 (6), 1543-1550 (1992)
                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
1 (bases 1 to 1803)
Spriggs,M.
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                                                                                                                                                           Submitted (28-JUL-1992) M. Spriggs, Immunex Res & Development Corporation, 51 University Street, Seattle WA 98101, USA
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H.sapiens mRNA for CD40 ligand.
X67878 S50586
X67878.1 GI:38411
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                                                                               Search completed: May 30, 2002, 02:49:21
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AL075432 Drosophil AL177093 Terracdon AL514085 AL514085 AQ373239 RPCIII-14 AL513979 AL513979 AL08993 Drosophil AL239086 Terracdon AL203738 SP.0084_A AL062806 Drosophil AL232112 Terracdon AL232112 Terracdon AL513817 AL513817 AL28200 Terracdon AL513817 AL513817 AL287681 Terracdon AL508200 Terracdon AL51757 HS_5184_B AL287681 Terracdon AL60999 Drosophil AL06199 Drosophil AL06199 Drosophil AL29763 Terracdon AL29763 Terracdon AL29763 Terracdon AL29763 Terracdon AL608989 Drosophil AL06088 Drosophil AL06088 Drosophil AL06088 Drosophil AL5146989 Drosophil AL5146989 Drosophil AL5146989 Drosophil AL5146989 Drosophil

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CNS011TX
A1557564
A1557564
CNS011TX
AL513817
CNS02BN3
CNS04BYW
CNS06BYL
CNS

ALIGNMENTS

	BF599437 492 bp mRNA linear EST 25-APR-2001 263218 MARC 3BOV Bos taurus CDNA 5', mRNA sequence.	BF599437		cow.	bos caurus Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;	Bovidae; Bovinae; Bos.	Smith, T.P.L., Grosse, W.M., Freking, B.A., Roberts, A.J., Stone, R.T.,	Casas, E., Wray, J.E., White, J., Cho, J., Fahrenkrug, S.C., Bennett	'G.L., Heaton,M.P., Laegreld,W.W., Rohrer,G.A., Chitko-McKown,C.G., Pertea,G., Holt,I., Karamycheva,S., Liang,F., Ouackenbush,J. and	Keele, J.W.	Sequence evaluation of four pooled-tissue normalized bovine cDNA	libraries and construction of a gene index for cattle	Genome Res. 11 (4), 626-630 (2001)	ZITROOTS	Contact: Smith TPL	USDA, ARS, US Meat Animal Research Center	PO Box 166, Clay Center, NE 68933-0166, USA	0001 000 0001	FdX: 404 /04 43VU	Email: Smitheemail:marc.usda.gov	single pass sequencing. Bases called and alt_trimmed with phred v0.980904.e. Vector identified by cross match with the -minscore 18	and -minmatch 12 options.	PCR PRimers	FORWARD: AGGAAACAGCTATGACCAT BACKWADD: CHHMHOOCACACAACAACAAC	DISTORAGE GILLLOCCHARGE	Seq primer: ATTAGGTGACACTATAG.	Location/Qualifiers	1.494 /organism="Bos taurus" /db_xref="taxon:9913"
RESULT 1 BF599437	LOCUS	ACCESSION	KEYWORDS	SOURCE	ORGANIZA	•	GUNAGAAAA					TITLE		JOURNAL	MEDLINE	COMMENT											-	FEATURES	e conce
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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gb_gss:* em_gss_hum

gb_est2: gb_htc:*

em_htc:* gb_est1:*

em_estba:*

EST:*

Database :

em_estin: em_estmn:

em_estov: em_estpl:

em_estro:

SUMMARIES

Query Match Length DB

Score

. 9

Description

BF599437 263218 MA
AM486605 75217 MAR
A1982044 par.pk007
AL534423 AL534423
AL063921 Drosophil
AL103735 Drosophil
AL069997 Drosophil
AL09999 Drosophil
AL00960 Drosophil
AL100660 Drosophil
AL514085 AL514085
AL105471 Drosophil
AL51373 Prosophil
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331.8 256.4 69.6

51.4 50.8 50.2

AL514657 AL514721

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rel: 402 762 4366
Fax: 402 762 4390
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Smith, T.P.L., Grosse, W.M., Freking, B.A., Roberts, A.J., Stone, R.T., Smith, T.P.L., Grosse, W.M., Freking, B.A., Fahrenkrug, S.C., Bennett Casas, E., Wray, J.E., White, J., Chorer, G.A., Chitko-McKown, C.G., G.L., Heaton, M.P., Lagereid, W.W., Rohrer, G.A., Chitko-McKown, C.G., G.L., Reaton, M.P., Lagereid, W.W., Edgereid, W.W., Rohrer, G.A., Chitko-McKown, C.G., G.L., Reaton, M.P., Lagereid, W.W., Roberts, G.A., Chitko-McKown, C.G., G.L., Reaton, M.P., Lagereid, W.W., Roberts, G.W., Chitko-McKown, C.G., C.C., C.G., C.C., C.G., C.
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Cetartiodactyla; Ruminantia; Pecora; Bovoldea
                                       Library made from pooled tissue from marrow, alveolar macrophage, ovary, fetal semitendonosus muscle, and fetal longissimus c. 104 a """
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Contact: Smith TPL
USDA, ARS, US Meat Animal Research Center
PPO Box 166, Clay Center, NE 68933-0166, USA
             /clone_lib="MARC 3BOV"
/tissue_type="pooled"
/lab_host="DH10B"
/note="Vector: pCMV SPORT6; Site
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75217 MARC 1BOV BOS taurus CDNA
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An expressed sequence tag database of T-cell-enriched activated chicken splenocytes: sequence analysis of 5251 clones Genomics 66 (2); 144-151 (2000)
Fax: 402 762 4390
Bmail: marc.usda gov
Bmail: marthemail.marc.usda gov
Single pass sequenching. Bases called and trimmed with phred
v0.580904 e. Vector identified by cross_match with the -minscore
and -minmatch 12 options:
PCR PRIMCTS
PORWARD: AGGAAACAGCTATGACCAT
                                                                                                                                                                                                                                                          /db_xref="taxon:9913"
/db_xref="taxon:9913"
/clone_lib="MARC 1B0V"
/tissue_type="pooled"
/lab_host="bH10B"
/note="vector: pCWV SPORT6; Site_1: xbaI; Site_2: XhoI; Library made from pooled tissue from lymph node, ovary, library made from plituitary."
fat, hypothalamus, and pituitary."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                241 TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 53 ATGATCGAAACATACAGTCAACCTTCTCCCGGCTCGGGGCCACTGGACCACTGTCAGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GAAGATTTTGTATTCATAAAAAAGCTAAAAGAGATGCAACAAAAGGAGAAGGATCTTTATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gallus gallus
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Archosauria; Aves; Neognathae; Galliformes; Phasianidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 398;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 256.4; DB 9
Pred. No. 3.7e-41;
0; Mismatches 46
                                                                                                                                   BACKWARD: GTTTTCCCAGTCACGACG
Plate: 34 row: F column: 23
Seq primer: ATTTAGGTGACTATAG.
                                                                                                                                                                                                                                           /organism="Bos taurus'
                                                                                                                                                                                                Location/Qualifiers
1. .398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GI:5885072
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Phasianinae; Gallus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        32.7%;
85.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AI982044.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
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560 CATTTATA 567
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CNS0039G/c
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DEFINITION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AL534423 TJ_FL013_FBrn1 Homo sapiens cDNA clone CSODF004YD24 5 prime, mRNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                          5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       614
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -GCGCCATTCACCCTCTATATTTATTGTACCTCCCCATGGAAGAGACCGGCTCCTGAT 115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       671
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAATGTGACTGATCCAAG 731
                                                                                                                                                                                                                                                                                                                                                                                                                          AGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAATCGGGAAGCTTCGAGTCA 554
                                                                                                                                                                                                                                                                                                                                                                                                                                                        26
                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - France
Email: segref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
_Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                         2 AGGGCTCTACTACATCTACTCACAAGTCAGCTTCTGCACCAAGGCGGGGGGTTCG----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       555 AGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGATTCGAGAATCTTACT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       615 CAGAGCTGCAAATACCCACAGTTCCGCCAAACCT---TGCGGGCAACAATCCATTCACTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    116 GAAGGGACTTGACACCCACAGCACCTCCACGCCTCTGTGAGCTCCAGTCCATCCGGGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
11. (bases 1 to 797)
11. (M. B., Gruber, C., Jessee, J. and Polayes, D.
Full.length cDNA libraries and normalization
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                         Length 638;
                                                                                                                                                                                                                                        /cell_type="Con A-activated splenic T cell"
/lab_host="E.coli TOP10 F'"
/note="Vector: pcDNA3"
/171 c 151 g 180 t 4 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   732 CCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAACTCT 781
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    236 AGCAGTGAACGTCAACCCTGGCAACACCTACTTTGGCATGTTCAAGCTGT 285
                                                                                                                                                                       // Ab zero = taxon:931
// Actione = pat.pk0072.c9.f"
// Clone = lib = "chicken activated T cell cDNA"
                                                                                                                                                                                                                                                                                                                                                         Score 69.6; DB 9; Length 6
Pred. No. 0.00038;
0; Mismatches 114; Indels
                                            40 Townsend Hall, Newark, DE 19717, USA Tel: 302 831-1345 Fax: 302-831-3411 Email: joan@UDel.Edu, www.chickest.udel.edu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /clone_lib="LTI_FL013_FBrn1"
                                                                                                                                                             /organism="Gallus gallus"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="CSODF004YD24"
                                                                                                                             Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AL534423.1 GI:12797916
             Molecular Endocrinology
University of Delaware
Contact: Joan Burnside
                                                                                                                                                                                                                                                                                                                                                         8.9%;
ilarity 57.6%;
Conservative
                                                                                                                                                                                                                          /sex="male"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Contact: Genoscope
                                                                                                              Seg primer: T7
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167; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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Best Local S
Matches 167
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DEFINITION
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AUTHORS
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COMMENT
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Chote="Corollary Corollary Corollary
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end of BAC #
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
/dev_stage="pooled tissue from post conception fetuses (20
desk, 24 week and 26 week)"
//sk host="hullot"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (02-JUN-1999) Genoscope - Centre National de Sequencage :
BP 191 91006 EVRY cedex - FRANCE (E-mail : segrefégenoscope.cns.fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Drosophila melanogaster genome survey sequence TET3 end of BAC # BACR08K10 of RPCI-98 library from Drosophila melanogaster (fruit AL063921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               140 ATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCATGAAGATTTTGTATTCATAA 199
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCTCAAATTGCGGCACATGTCATAA 379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TGAGCAACAACTTGGTAACCCTGGAAATGGGAAACAGCTGACCGTTAAAAGACAAGGAC 499
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         240 AGAAGAAAAAAAAAAATTWAAAAAAGAAAAATTAAATGAAAAAAGGAAAAATAA 299
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         200 AAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCCTTGCTGAACTGTGAGGAGA 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  300 AGGAAAATRTAAAAAAAAAAAAAAGTAAATTWAWTTWAWRRAGGATAAAAATAAAAAAA 359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAGAAGAGAAAAAAG 319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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Drosophila melanogaster

Bukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;

Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;

Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.

1 (bases 1 to 1101)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     225;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 54; DB 9;
Pred. No. 0.45;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                35; Mismatches
                                                                week, 24 week and
/lab_host="DH10B"
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Matches 168; Conservative
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us-08-982-272-20.rst

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scomorpha; Ephydroidea; Drosophilidae; Drosophila.
(bases 1 to 1043)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6.68;
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                                                                                                                                                                                                                                                                                                                                                                                                                                               96 c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                        Direct Submission
Submitted (23-JUL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Muscomorpha;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Genoscope
                                                                                                                                                                                                                                                                                  pBeloBAC11
                                                            Senoscope
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Direct
                                                                                                                                                                                                                                                                                                                                                                                                                                                 277
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local S:
Matches 107
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                                                                        The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further information please see http://www.fruitfly.org The BDGP brosophila melanogaster BAC library was prepared by Kazutoyo Osoegawa and Aaron Mammoser BAC library was prepared by Kazutoyo Osoegawa and Aaron Mammoser an Pieter de Jong's laboratory in the Department of Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPCI-98 and was constructed by partial ECORI digestion of Drosophila DNA provided by the BDGP from the isogenic strain y2, on by the same strain used for the BDGP's pl and BST libraries. A more detailed description of the library and how to order individual BAC clones, the entire library, or filters for hybridization from the BACPAC Resource Center can be found at http://Dacpac.med.buffalo.edu/drosophila_bac.htm.
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                                               sequence was carried out as part of a
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BACN11G11 of DrosBAC library from Drosophila melanogaster (fruit
                                                  Determination of this BAC-end sequence was carried out as part of collaboration with the Berkeley Drosophila Genome Project (BDGP).
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survey sequence T7 end of BAC
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Neoptera, Endopterygota, Diptera, Brachycera,
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tive 199; Mismatches 146;
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/clone_lib="RPCI-98"
/clone="BACR08K10"
                                   : www.genoscope.cns.fr)
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of BAC #
                                             - Web : www.genoscope.cns.fr)
Determination of this BAC-end sequence was carried out as part of a Determination of this BAC-end sequence was carried out as part of a Determination of this BAC-end sequence brosphila Genome Project (EDGP) - http://www.edgp.ebi.ac.uk -. This Drosophila melanogaster BAC library (Dros BAC) was made by Alain Billaud at CEPH (Centre C'Etude du Polymorphisme.Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton and Genevieve Payan. It has been constructed in the vector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CNSOO6MT BACR14K09 of RPCI-98 Library from Drosophila melanogaster genome survey sequence TET3 end of BAC #BACR14K09 of RPCI-98 Library from Drosophila melanogaster (fruit
                      - FRANCE (E-mail : segref@genoscope.cns.fr
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s; Ephydroidea; Drosophilidae; Drosophila.
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                                                                                                                                                                                                                                                                                                                                         /plasmid="pBeloBAC11"
/db_xref="taxon:7227"
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/note="end : T7"
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                          Determination of this BAC-end sequence was carried out as part of a collaboration with the Barkeley brosophila Genome Project (BDGP). The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further information please see http://www.fruitfly.org The BDGP Drosophila melanogaster BAC library was prepared by Kazutcyo Osoegawa and Aaron Mammoser in Pieter de Jong's laboratory in the Department of Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPCI-98 and was constructed by partial ECORI digestion of Drosophila DNA provided by the BDGP from the isogenic strain y2; cn bw sp, the same strain used for the BDGP's
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                      segref@genoscope.cns.fr
Submitted (02-JUN-1999) Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - FRANCE (E-mail : segref@genoscope.cns.fr
                                                                                                                                                                                                                                                      isogenic strain y2; on bw sp, the same strain used for the BDGP's Pland EST libraries. A more detailed description of the library and how to order individual BAC clones, the entire library, or filters for hybridization from the BACPAC Resource Center can be
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                  found at http://bacpac.med.buffalo.edu/drosophila_bac.htm
Location/Qualifiers
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/db_xref="Laxon:7227"
/clone_lib="RPCI-98"
/clone="BACR14K09"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 226
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 12;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 50.8;
Pred. No. 1.
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33.9%; Pred
73;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               307
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Best Local S
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                                                                                                                                                                                                                                                                                                                                                   collaboration with the Berkeley brosophila Genome Project (BDGP).

The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further information please see http://www.fruitfly.org The BDGP Drosophila melanogaster BAC library was prepared by Kazutoyo Osoegawa and Aaron Mammoser in Pieter de Jong's laboratory in the Department of Cancer Genetics at the Roswell Park Cancer Institute in Buffalo, NY. The library is named RPCI-98 and was constructed by partial Eogenic strain y2; on bw sp., the same strain used for the BDGP's pland and EST libraries. A more detailed description of the library
                                                                                                                                                          Genoscope.

Direct Submission
Submitted (02-JUN-1999) Genoscope - Centre National de Sequencage :
BP 191 91006 EVRY cedex - FRANCE (E-mail : segref@genoscope.cns.fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      filters for hybridization from the BACPAC Resource Center can be found at http://bacpac.med.buffalo.edu/drosophila_bac.htm.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGGGCTGAAAAAGGATACTACACCA 439
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      881 ARARARRARAKAARARARGARRRRGRRRRRGRAGRRRARGAGGRAGRRGRGGRAGRRARARR 822
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAAAGCTAAAGAGTGCAACAAAGGAGAAGGATCTTTATCCTTGCTGAACTGTGAGGAGA 259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               260 TGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAAAGAAGAAAAAG 319
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       641 RERARGARRERRGRRERRERRARGRRERRAGRGRREGRRAGGARRGARGARRAGRRERRARA 582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          and how to order individual BAC clones, the entire library, or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   140 ATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCATGAAGATTTTGTATTCATAA 199
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Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
1 (bases 1 to 997).
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                                                                                                                                                                                                                                                                                           - Web : www.genoscope.cns.fr)
Determination of this BAC-end sequence was carried out as
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/db_xref="taxon:7227"
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Pred. No. 2.4;
}; Mismatches 156;
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/clone="BACR12K22"
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Drosophila melanogaster
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Genoscope
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Best Local S
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                                                                                                                                  ORGANISM
                            DEFINITION
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TITLE
JOURNAL
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                                                                                    VERSION
KEYWORDS
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                                                                                                                                                                                                      web : www.genoscope.cns.fr)
Determination of this BAC-end sequence was carried out as part of scollaboration with the European Drosophila Genome Project (EDGP) thtp://www.edgp.ebi.ac.uk - This Drosophila melanogaster BAC library (Dros BAC) was made by Alain Billaud at CEPH (Centre d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton and Genevieve Payan. It has been constructed in the vector
                                                                                                                                                                              Submitted (23-JUL-1999) Genoscope - Centre National de Sequencagé
BP 191 91006 EVRY cedex - FRANCE (E-mail : segref@genoscope.cns.fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         675 MACACVMSMWCASACAMCACACACACACAMCMMTMACCMCCSSMASSSNAASASYVTVV 734
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Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
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                                                                                                                                                                                                                                                                                                                                                                      /organism="Drosophila melanogaster"
/plasmid="pBeloBAC11"
/db_xref="taxon:7227"
/clone_lib="DrosBAC"
/clone="BACNOSCOS"
/note="end: T7"
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27.1%; Pred. No. 6.9;
tive 88; Mismatches 227;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     287
         survey sequence.
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                                                                                Drosophila melanogaster
                                     AL099997.1 GI:5611608
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           fly), genomic
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                                                                   fruit
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Best Local S
Matches 117
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                         ACCESSION
VERSION
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                                                                                                                                                                         TITLE
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RESULT

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p DNA linear GSS 26-JUL-1999
survey sequence T7 end of BAC
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- Web : www.genoscope.cns.fr)

- Determination of this BAC-end sequence was carried out as part of collaboration with the European Drosophila Genome Project (EDGP) - http://www.edgp.ebi.ac.uk - This Drosophila melanogaster BAC library (Dros BAC) was made by Alain Billand at CEPH (Centre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton and Genevieve Payan. It has been constructed in the vector
                             planogaster genome survey sequence T7 end of BAC DrosBAC library from Drosophila melanogaster (fruit
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Muscomorpha, Ephydroidea, Drosophilidae, Drosophila.
1 (bases 1 to 799)
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/plasmid="pBeloBAC11"
/db_xref="taxon:7227"
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Pred. No. 9.1;
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Eukaryota; Metazoa; Arthropoda;
              799 bp
                                                                      fly), genomic survey sequence.
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                                       Drosophila melanogaster
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/note="end : T7
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                                                                                                                                            Direct Submission
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                     fruit fly:
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/note="Vector: pcMvSPORT 6; Site_1: NotI; 1st strand cDNA
was primed with a NotI-oligo(dT) primer. Five prime end
enriched, double-stranded cDNA was digested with Not I and
cloned into the Not I and Eco RV sites of the pcMvSpORT 6
vector Library was normalized. Library was constructed by
Life Technologies. Contact: Feng Liang Life Technologies,
a division of Invitrogen 9800 Medical Center Drive
Rockville, Maryland 20850, USA Fax: (1) 301 610 8371.
Email: fliang@lifetech.com URL:
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                     458 bp mRNA linear EST 13-FEB-2001
LTI_NFL006_PL2 Homo sapiens cDNA clone CL0BA0042E10 3
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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AL105471
AL105471.1 GI:5617485
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Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - France
Bmail: segref@genoscope.cns.fr, Web : www.genoscope.cns.fr.
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       6.0%; Score 46.8; DB 9; Length 458; 42.5%; Pred. No. 14; Live 14; Mismatches 144; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     49 others
                                                                                                                                                                                       1 (bases 1 to 458)
Lil,W.B., Gruber,C., Jessee,J. and Polayes,D.
Unll-length cDNA libraries and normalization
Unpublished (2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   //fulllength.invitrogen.com
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                                                                                                                                                                                                                                                                                                                                                                 /organism="Homo sapiens"
/db_xref="taxon:9606"
/clone="CL0BA0042E10"
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                                                                                         AL514085.1 GI:12777579
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prime, mRNA sequence.
AL514085
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Matches 117; Conservative
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                                                                                                                                            Homo sapiens
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                                                                                                                           human.
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CNS0127J 858 bp DNA linear GSS 26-JUL-1999
Drosophila melanogaster genome survey sequence T7 end of BAC
BACN07D07 of DrosBAC library from Drosophila melanogaster (fruit
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                                                                                                                                                                                                                                            Web: www.genoscope.cns.fr)
Determination of this BAC-end sequence was carried out as part of a collaboration with the European Drosophila Genome Project (EDGP) - http://www.edgp.ebi.ac.uk - . This Drosophila melanogaster BAC library (Dros BAC) was made by Alain Billaud at CEPH (Centre G'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton and Genevieve Payan. It has been constructed in the vector
                                                                                                                                                                                                                           segref@genoscope.cns.fr
                                                                                                                                                                                              de Sequencage
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Drosophila melanogaster

Drosophila melanogaster

Enkaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;

Enkaryota; Metazoa; Arthropoda; Diptera; Brachycera;
                                     Insecta;
ruerygota; Neoptera; Endopterygota; Tracheata; Hexapoda; Ins. Muscomorpha; Ephydroidea; Drosophilidae; Drosophila. 1 (bases 1 to 1064)
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Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
1 (bases 1 to 858)
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                                                                                                                                                                                        Submitted (23-JUL-1999) Genoscope - Centre National BP 191 91006 EVRY cedex - FRANCE (E-mail : segref@ge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                169 others
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Indels
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Similarity 32.4%; Pred. No. 13;
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/organism="Drosophila
/plasmid="pBeloBAC11"
/db_xref="taxon:7227"
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/clone="BACN14C18"
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AUTHORS
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us-08-982-272-20.rst

TITLE JOURNAL

COMMENT

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ALS13719 LTI_NFL006_PL2 Homo saplens CDNA clone CL0BA0072H01 3
                                                Web: www.genoscope.cns.fr)
Determination of this BAC-end sequence was carried out as part of a collaboration with the European Drosophila Genome Project (EDGP) - http://www.edgp.ebi.ac.uk - fnis Drosophila melanogaster BAC library (Dros BAC) was made by Alain Billaud at CEPH (Centre & Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton and Genevieve Payan. It has been constructed in the vector
                  Submitted (23-JUL-1999) Genoscope – Centre National de Sequencage
BP 191 91006 EVRY cedex – FRANCE (E-mail : seqref@genoscope.cns.fr
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1 (bases 1 to 329)
Li,W B., Gruber,C., Jessee,J. and Polayes,D.
Full-length cDNA libraries and normalization
Unpublished (2001).
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/clone="BACN37P10"
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Eukaryota; Metazoa;
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Direct Submission
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         Direct Submission Submitted (23-Jul-1999) Genoscope - Centre National de Sequencage: Submitted (23-Jul-1999) Genoscope - Centre National de Sequencage: PP 191 91006 EVRY cedex - FRANCE (E-mail: seqref@qenoscope.cns.fr - Web: www.genoscope.cns.fr)

Determination of this BAC-end sequence was carried out as part of a collaboration with the Buropean Drosophila Genome Project (EDGP) - http://www.edgp.ebi.ac.uk - This Drosophila melanogaster BAC thibrary (Dros BAC) was made by Alain Billaud at GEPH (Centre d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton and Genevieve Payan. It has been constructed in the vector
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Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila
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/clone="BACN07D07"
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Best Local Similarity 37.09
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14

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BASE COUNT ORIGIN

FEATURES

39

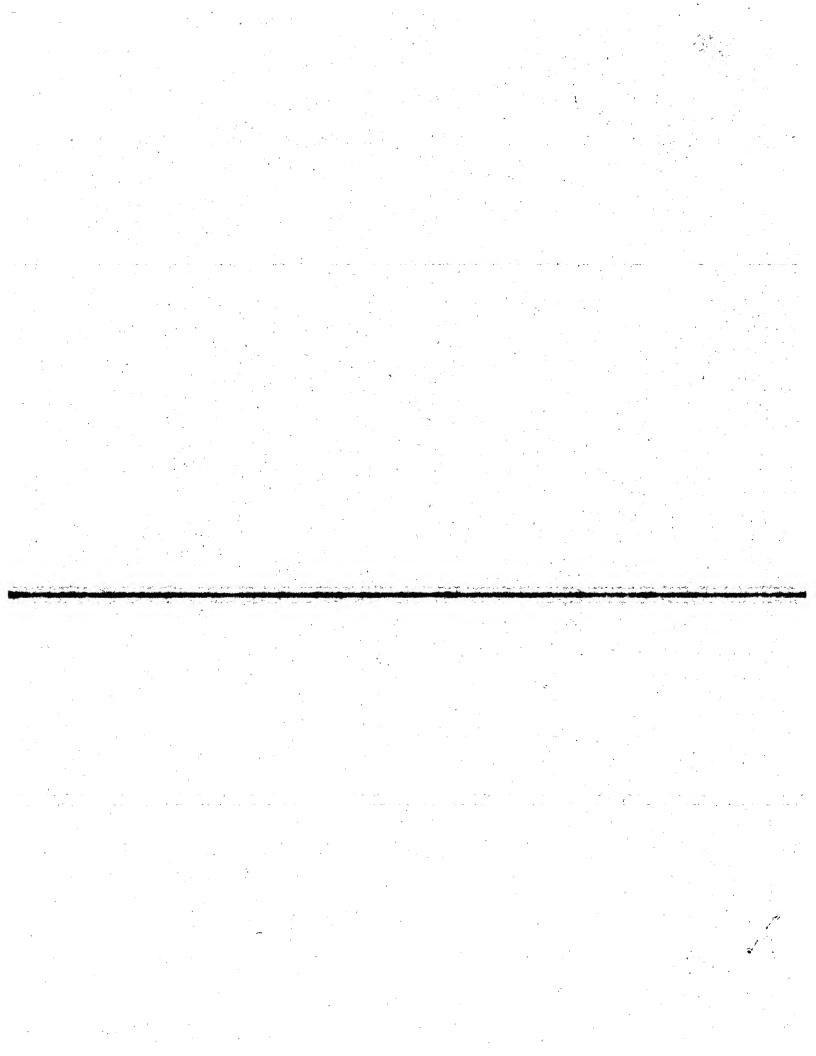
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/dbysitish - Taxon: Saptens / dbysitish - Taxon: Saptens / dbysitish - Taxon: 9606 * / clone="CLOBA0072H01" / clone="Lib="Lil"LTIL" INFLO06_PL2" / tlssue_lype="Polacenta" / tlssue_type="Polacenta" / tlssue_type="Polacenta" / note="Vector: pcMvSPORT 6; Site_l: NotI; lst strand cDNA was primed with a NotI-oligo(dT) primer. Five prime end enriched, double-stranded cDNA was digested with Not I and cloned into the Not I and Eco RV sites of the pcMvSPORT 6 vector. Library was normalized. Library was constructed by Life Technologies. Contact: Feng Liang Life Technologies, a division of Invitrogen 8000 Medical Center Drive Rockville, Maryland 20850, USA Fax: (1) 301 610 8371 Email: fliang@lifetech.com URL: http://fulllength.invitrogen.com R7 3 9 60 t 67 others
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                       Genoscope - Centre National de Sequencage
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Email: seqrefégenoscope.cns.fr, Web : www.genoscope.cns.fr.
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                                                                                                                                                  /organism="Homo sapiens"
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1 ATGATCGAAACATACAACCA.....TTGGCTTACTCAAACTCTGA 783
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

/SIDS1/gcgdata/hold-geneseq/genesegn-embl/NA2002.DAT:*

/SIDS1/gcgdata/hold-geneseq/genesegn-embl/NA2001A.DAT /SIDS1/gcgdata/hold-geneseg/genesegn-emb1/NA2001B.DAT

Description	Exemplary nucleoti Human CD40-L type CD40 ligand gene u CD40 ligand coding CD40-L DNA. Homo CD40 ligand gene. Human CD40 ligand CDNA of CD40L, a n Human CD40 ligand
SUMMARIES	AAV42184 AAV58359 AAV12852 AAQ41506 AAQ67123 AAT07763 AAT07763
OB	119 119 119 118 119
8 Query Score Match Length DB I	783 786 786 786 786 840 840 840 840
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υ	. 32	599.2	76.5	780	21	AAZ55543	٠	Feline CD154 cDNA	
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-		280	74.1	780	21	AAZ55536 '		CD154	
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altering the immunoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface of the cells. Vectors containing the
                                                                                                             The present sequence represents an exemplary nucleotide sequence, where nucelotides encoding human domain III have been replaced with nucleotide encoding mouse domain III. The sequence is used to exemplify the method of the invention. The specification describes a method for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACCGTTAAAAGACAAGGACTCTATTÄTATCTATGCCCCAAGTCACCTTCTGTTCCAATCGG 540
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autoimmune disorders such as rheumatoid arthritis. They can also be
used for vaccination to produce immunity against a virus cell,
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Vectors containing accessory molecule ligand genes - used for altering immunoreactivity of cells, particularly for treatment of neoplasia or autoimmune disorders, e.g. rheumatoid arthritis
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                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Score 783; DB 19;
llarity 100.0%; Pred. No. 7e-207;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                           protein, fungus or neoplasia.
                                                                               Disclosure; Page 116; 167pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         t= human_CD40-L
"nucleotides 148-783 code for the
extracellular region (amino acids 50-261)"
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                                                                                                                                                                                                                                                                                                                  Leucine zipper; trimerisation; trimeric CD40-L; fusion protein; hetero-oligomer; homo-oligomer; type II transmembrane protein; soluble CD40-L; tumour necrosis factor family; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            cells to express a fusion protein comprising a leucine zipper domain and a heterologous mammalian protein
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                                                                                                                                                                                                                                                                                      Human CD40-L type II transmembrane protein coding sequence.
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Pred. No. 2.6e-191;
); Mismatches 27;
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                                                                                                                                                                                        AAQ63959 standard; cDNA to mRNA; 786
                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
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 atgatcgaaacatacaaccaaacttctccccgatctgcggccactggactgcccatcagc
                                   ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGC
                                                                                                     GAAGATTTTGTATTCATAAAAAGCTAAAGAGGTGCAACAAAGGAGGATCTTTATCC
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                                                                                                                                                                                                                                                                                                                                                                                                               accessory molecule ligand, AML, gene therapy; treatment, autoimmune disorder, rheumatoid arthritis; vaccine; ss.
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of the cells. Vectors containing the AML genes can be used in gene therapy for treating neoplasia or autoimmune disorders such as rheumatoid arthritis. They can also be used for vaccination to produce immunity against a virus cell, bacteria, protein, fungus or neoplasia.
                                                                                                                                                                                                                                                                                                                                                                              used to exemplify the method of the invention. The specification describes a method for altering the immunoreactivity of human cells which comprises introducing a gene encoding an accessory molecule ligand (AML) into the cells so that the AML is expressed on the surface
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Score 726.8; DB 19;
; Pred, No. 2.6e-191;
. 0; Mismatches 27;
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96.2%;
                                       97WO-US22740
                                                                      97US-0982272
96US-0032145
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Best Local Similarity 96.2
Matches 756; Conservative
                                                                                                                              (REGC ) UNIV CALIFORNIA
                                                                                                                                                                     Kipps TJ,
                                                                                                                                                                                                         WPI; 1998-348521/30
                                     08-DEC-1997;
                                                                        01-DEC-1997;
                                                                                          09-DEC-1996;
                                                                                                                                                                   Cantwell M,
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18-JUN-1998
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The encoded protein can be used in a fusion protein produced using the method of the invention. The method is for preparing soluble oligomeric protein by culturing a host cell transfected with a vector for the fusion protein. The soluble oligomeric proteins comprise a leucine zipper fused to the N terminus of the extracellular region of a heterologus mammalian type II transmembrane protein or to the C terminus of the extracellular region of a heterologus mammalian type II transmembrane
                                                                                                                                                                                                                                                                                                                          heterologous mammalian type II transmembrane protein; activated T cell; heterologous mammalian type I transmembrane protein; activated T cell; cheterologous mammalian type I transmembrane protein; antibody production; CD40-L; B-cell proliferation; CD27-L; lymphocyte antigen; ss.
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          cgggaagettcgagtcaagetccatttatagccageetetgeetaaagteeeeggtaga 600
                               CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
sequence is the coding sequence for the human CD40 ligand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          comprising leucine zipper
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to extracellular region of transmembrane protein
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91US-0783707.
91US-0805723.
92US-0969703.
                                                                                                                                                                                                                                                                                                           CD40 ligand coding sequence.
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of AAM41171 or AAM41172, optionally with conservative amino acid substitutions, provided that the peptide trimerises in solution. A soluble fusion protein comprising the leading tipper of AAM41171 linked to the extracellular region of CD40. (a type II transmembrane protein antigen CD40) stimulates B-cell and acts as a ligand for the B-cell antigen CD40) stimulates B-cell proliferation and antibody production in a similar manner to membrane-bound CD40.L. A soluble fusion protein comprising the leucine zipper of AAM41172 linked to the extracellular region of CD27-L (a type II transmembrane protein that binds to the ilymphocyte antigen CD27) inhibits binding of CD27c (a fusion protein comprising the extracellular region of CD27 and a human IgG1 Fc region) to BBV-transformed B cells expressing CD27-L.
a peptide comprising at least part
               protein, where
of AAW41171 or
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Sequence 786 BP; 250 A; 168 C; 168 G; 200 T; 0 other;

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GTGACTCATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 777
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                                                                                                                                                                                                                                                                                                     gotgaaaaaggatactacaccatgagcaacattggtaaccctggaaatgggaaacag
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                                                                                 CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT
                                                                                                                                                       1 ATGATCGAAACATACAACCTACTCCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
                                                     1 atgatcgaaacatacaaaccaaacttctccccgatctgcggccactggactgccatcagc
                                                                        ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA
                                                                                                                                               GAAGATTTTGTATTCATAAAAAAGCTAAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
                                             Length 786;
                   Indels
 DB 19;
Score 726.8; DB 19;
Pred. No. 2.6e-191;
; Mismatches 27;
    Score
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  92.8%;
Local Similarity 96.2%;
hes 756; Conservative C
    Query Match
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                                                                 TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA
                                                                                                                                                            AACAAAGAAGA - - - GAAAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                                                                                                                                         GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG
                                                                                                                                                                                                                                                                                                                                CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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                                                    GAAGATTTTGTATTCATAAAAAAGCTAAAGAGATGCAACAAAGGAGAGGATCTTTATCC
                                                                                                                                                                                                                  CAAATTGCGGCACATGTCATAAGTGAGGCCCAGCAGTAAAACAACATCTGTGTTACAGTGG
                                                                                                                                                                                                                                                                                                                                             /*tag= a
/product= CD40 ligand
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CD40-L is a type II membrane polypeptide which has an extracellular region at its C-terminus, a transmembrane region and an intracellular region at its N-terminus. A soluble form of CD40-L lacks the transmembrane domain. CD40-L activity is mediated by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           binding with CD40 an dinduces B cell proliferation and induction of antibody secretion, including IgE. Membrane bound CD40-L acts as a CD40 agonist and soluble CD40-L acts as a CD40 antagonist. CD40-L can be used in a binding assay to detect cells expressing CD40.
                                                                                                                                                                                          transmembrane; region; intracellular; soluble; activity; B cell; proliferation; induction; antibody; secretion; IgE; agonist; antagonist; binding assay; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New cytokine CD40-L as CD40 agonist and antagonist - is used for treating allergies, lupus, rheumatoid arthritis, graft-versus-host disease and insulin-dependent diabetes mellitus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This sequence encodes a human CD40-L polypeptide which binds to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 726.8; DB 14;
Pred. No. 2.7e-191;
0; Mismatches 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Spriggs MK;
                                                                                                                                                                                                                                                                                                 Location/Qualifiers
46..831
/*tag= a
                                                                                         AAQ41506 standard; DNA; 840 BP
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Best Local Similarity 96.2%;
Matches 756; Conservative
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91US-0805723
                                                                                                                                             (first entry)
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P-PSDB; AAR36701.
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  783
                 Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                          25-OCT-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-DEC-1991;
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778 CTCTGA
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                                                                                                                                                                        CD40-L DNA
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                                                                                                                                                                                                                                                                         this gene were identified by the method of the invention. The method comprises isolating DNA from an individual and selectively amplifing the isolated DNA derived from the CD40 ligand gene. The amplification product is then analysed to determine if there is a mutation present and determining if a protein expressed from the ligand gene will bind CD40. The detection of mutations in the CD40 ligand gene allows subsequent treatment of a syndrome resulting in elevated levels of serum igh and diminished levels of other Igherory, and the mutation in the CD40 ligand gene. ie. X-linked
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GAAGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
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                                                                                                                                                                                                                                                              Mutations within
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 840;
                                                                                                                                                                                            - comprises
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                                                                                                                                                                                          mutation in CD 40 ligand gene - colleic acid, and mutational analysis
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 840 BP; 266 A; 185 C; 175 G; 214 T; 0 other;
                                                                                                     Renshaw BR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 726.8; DB 15;
Pred. No. 2.7e-191;
0; Mismatches 27;
                                                                                                                                                                                                                                                                sequence represents the CD40 ligand gene.
                                                                                                         Fanslow WC,
                                                                                                                                                                                                                                       Disclosure; Page 22-24; 38pp; English.
                                                                                                                                                                                            Method for detecting mutation I amplification of nucleic acid,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 92.8%;
Best Local Similarity 96.2%;
Matches 756; Conservative
       94WO-US00786
                                   93US-0009258
                                                 94US-0184422
                                                                                                         Davison BL,
                                                                                                                        Widmer MB;
                                                                             IMMV ) IMMUNEX CORP.
                                                                                                                                                  WPI; 1994-264109/32.
                                                                                                                                                                                                                                                                                                                                                                                                                      hyperigM syndrome
                                                                                                                                                                  P-PSDB; AAR57469
                                                                                                            RJ,
                                   22-JAN-1993;
20-JAN-1994;
       21-JAN-1994;
                                                                                                                      Spriggs MK,
                                                                                                            Armitage
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The nucleotide sequence given in AAT05763 encodes a human high-density.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            High density membrane bound CD40 ligand - for stimulating the proliferation of B cells in vitro or in vivo, partic. for producing differentiated cells
                                                                                                                                                                                                                                                                                     GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
                                           586 egggaagettegagteaagetecatttatageeageetetgeetaaagteeeeggtaga
                                                                                                                 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
                                                                                                                                                                                                                TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAAGTCCCCCGGTAGA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       membrane-bound CD40 ligand; B-lymphocyte; B-cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       High density membrane-bound CD40 ligand; B-lymphc differentiation; proliferation; baculovirus; Spod Sf9; insect cell culture; tumour necrosis factor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (BOEH ) BOEHRINGER INGELHEIM PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
22.807
/*tag= a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human CD40 ligand DNA.
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P-PSDB; AAR85486
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ctctga 831
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                                                             61 ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
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                                       1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGCCATCAGC
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                                                                                                                                         gaagattttgtattcatgaaaacgatacagagatgcaacacaggagaaagatccttatcc
                                                                                                                                                        TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA
                                                                                                                                                                                      AACAAAGAAGA - - - GAAAAAAAAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                                                                                    CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTTACAGTGG
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                3;
Length 840;
DB 16;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               cDNA of CD40L, a novel cytokine ligand for CD40
Score 726.8; DB 1 Pred. No. 2.7e-191
               0; Mismatches
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92.8%;
96.2%;
                Conservative
       Similarity
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ctctga 807
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ligand for CD40 called CD40L. CD40L is a type II membrane polypeptide, which is expressed by activated CD4+ T calls, and causes B cell proliferation and induction of antibodies, which in turn bind to CD40-expressing cells. This inhibits binding of soluble CD40 to its ligand CD40L. The monoclonal antibody against CD40L is used to inhibit proliferation of neoplastic cells, and is particularly useful in treating B-cell lymphoma (e.g. where induced after transplants or in other cases of immune deficiency such as AIDS), and also metanoma or carcinoma. Since the monoclonal antibodies inhibit neoplastic cells directly, they may not need to be coupled to a toxin or radioisotope, avoiding toxic effects on
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    present sequence represents the cDNA sequence of a novel
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Treating or preventing neoplastic disease associated with CD40-expressing cells - particularly B-cell lymphoma, by administration of CD40-binding protein, preferably antibody solubbe CD40-ligand
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Pred. No. 2.7e-191
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                                                                                          Location/Qualifiers
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(USSH ) US DEPT HEALTH & HUMAN
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96.2%;
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Matches 756; Conservative
  SS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fanslow
melanoma; carcinoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1997-502273/46.
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                                             Homo sapiens
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GAAGATTTTGTATTCATAAAAAAGCTAAAAGAGATGCAACAAAGGAGGAGGATCTTTATCC

181 226 AACAAAGAAGA - - - GAAAAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT

301

TTGCTGAACTGTGAGGAGATGAGGAAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA

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GCTGAAAAAGGATACTACACCATGAGCAACATTGGTAACCCTGGAAAATGGGAAACAG 477
                                                                                                                                                                                                                                                                                                                                                                          membrane glycoprotein; cell proliferation; differentiation;
                                                               TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                         CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAAACAACATCTGTGTTACAGTGG
                                                                                      CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                     CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                                                                                                                                                                                                                                                                                                                                                    activated T cell;
                                                                                                                                                                                                                                                                                                                                                                     Human; CD40 ligand; TNF receptor family;
                                                                                                                                                                                                                                                                                                                                                     Human CD40 ligand encoding DNA sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                           "CD40 ligand"
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P-PSDB; AAW71751.
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28-APR-1994;
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The present sequence encodes human CD40 ligand which is used in the method of the invention. The method has been developed for proliferating B cells to increase their number at least 100-fold. The method comprises: (a) providing high density, membrane bound CD40 ligand, and coulture results in a proliferation in the presence of this ligand. The culture results in a proliferation in the number of B cells of at least indoced to differentiate into antibody-producing cells in the presence of one or more cytokines. The method can be used for stimulating B-cell proliferation and activation is suppressed. Eight rounds of division over six days can be achleved, corresponding to a 256-fold increase in cell numbers, which is a vast increase compared to previous
by incubating them
    ation of B cells in culture
membrane-bound CD40 ligand
                                                                      Example 1; Fig 1; 37pp; English
         proliferation
                                in the presence of
                                                                                                                                                                                                                                                                                                                                                                                                  culturing methods.
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Length 840; Sequence 840 BP; 263 A; 182 C; 181 G; 214 T; 0 other; Score 726.8; DB 19; Pred. No. 2.7e-191; 92.8%; 96.2%; Query Match Best Local S

CAAICCATICACIIGGGAGGAGTAIITGAAIIGCAACCAGGIGCIICGGIGIITGICAAI 717, 180 201 300 357 381 441 477 537 621 657 321 501 141 261 TITGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA ATGAAAATTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA CITITITGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT GCTCAAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAAGA CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT 22 atgatcgaaacatacaaccaaacttctccccgatctgcggccactggactgccatcagc GAAGATTTTGTATTCATAAAAAGCTAAAGAATGCAACAAAGGAGAAGGATCTTTATCC ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC. 0; Mismatches Best Local Similarity 96.2 Matches 756; Conservative 442 502 538 562 418 478 298 622 658 61 142 241 181 g δ g ò g Ωp ď qq Ω

g δλ q Q δλ

537 585 597 645 657 705 717 765 777

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us-08-982-272-20.rng

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.gp39; CD40 ligand; osteoblast cell death; apoptosis; bone loss;
osteoporosis; osteoonecrosis; inflammatory arthritis; esterogen loss;
ovariectomy; historectomy; lupus nephritis; Takayasu's arteritis;
Wegener's granulomatosis; nephritis; myositis; scheroderma;
                                             346 aacaaagaggagacgaagaaagaaacagctttgaaatgcaaaaaggtgatcagaatcct
                                                                                                                                                                                                                                                                                                                                                                                                                                               CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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                         GAAGATTTTGTATTCATAAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
                                                                                                         241 TTGCTGAACTGTGAGGAGGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA
                                                                                                                                                                                           301 AACAAAGAAGA---GAAAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                                                                                                                                                CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
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                                                                                                                                                                                                                   Nucleotide sequence of human gp39 protein, a CD40 ligand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                thrombocytopenia; asthma; lung disease; cancer; ss
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This sequence encodes the human CD40 receptor ligand (CD40-L). The invention relates to anti-human CD40-L monoclonal antibodies M90 secreted by hybridoma hCD40L-M90 (ATCC HB-12055) and M91 secreted by hybridoma hCD40L-M91 (ATCC HB-12056). M90 and M91 inhibit CD40-L binding to CD40 and the ability of trimeric CD40-L and anti-Immunoglobulin M to induce
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                                                                                                                                                                                                                                                                                                                                                                                                                                         CD40-L; CD40 receptor ligand; anti-human CD40-L monoclonal antibody; binding inhibitor; trimeric CD40-L; anti-Immunoglobulin M; peripheral blood B cell; proliferation inhibitor; ss.
                                                              1 ATGATCGAAACATACAACCTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
  caatccattcacttgggaggagtatttgaattgcaaccaggtgcttcggtgtttgtcaat
                                         GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
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Pred. No. 2.7e-191;
0; Mismatches 27;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Anti-human CD40-Ligand monoclonal antibodies
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                                                                                                                                                                                                                                                                         AAZ27525 standard; cDNA; 840 BP
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P-PSDB; AAY39938.
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disease, nephrotic/nephrific syndrome, or cancer. They may also be to treat or prevent bone loss in a subject undergoing, or scheduled an organ or bone marrow transplant.
                                                                                                                                                                                                                                                                                                                                                                                                         sent sequence encodes a 9939 protein. It is a CD40 ligand. Jands are used for reducing osteoblast cell death or apoptosis, treating or preventing bone loss in animals, preferably humans, of, or undergoing, bone loss. The bone loss is associated with rosis, osteonecrosis, inflammatory arthritis, post menopausal.
                                                                                                                                                                                                                                                                CD40 agonist containing composition, used to reduce bone cell death or apoptosis associated with osteoporosis, osteonecrosis and inflammatory arthritis -
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Pred. No. 2.7e-191;
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ilarity 96.2%;
Conservative (
2000WO-US23276
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24-AUG-2000;
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cD40 engagement on T cells can be used to induce T cell receptor rearrangement and enhance T cell affinity for a particular antigen. CD40 engagement can be bought about by contacting CD40 with a CD40-binding agent; e.g. human cD40 ligand: The CD40-binding agents can also be used in methods for promoting T cell maturation, inhibiting T cell receptor rearrangement, inhibiting environmental stress-induced cell death, altering the specificity of a T cell towards an antigen, inducing T cell reactivity towards an antigen or enhancing environmental stress-induced cell death (all claimed). T cell affinity maturation towards a specific
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  cell death inhibition; stress-induced; immunosuppressive; anti-thyroid; anti-inflammatory; anti-diabetic; anti-rheumatic; anti-anaemic; opthalmological; anti-psoriatic; nephrotophic; hepatotropic; virucide; dermatological; cytostatic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CD40; T cell; T cell receptor; rearrangement; maturation;
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                                                                                                                                                                                      GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
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CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
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                                                                                                                                                                                                                                                                                                                                                              AAA51745 standard; cDNA; 1816 BP
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AAS86571 standard; cDNA; 1816 BP.

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            disease, which includes rheumatoid arthritis, uveitis, insulln-dependent diabetes mellitus, haemolytic anaemias, rheumatic fever, Crohn's disease, Guillain-Barre syndrome, psoribasis, thyroiditis, Grave's disease, myasthenia gravis, glomerulonephritis, autoimmune hepatitis or systemic lupus erythematosus. Inducting environmental stress induced T cell death is carried out in a cancerous T cell or a self-reactive T cell where the environmental stress is a chemotherapeutic agent (claimed).
  self-antigen in an autoimmune
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96.2%; Pred. No. 3.6e-191;
ive 0; Mismatches 27;
can be inhibited, especiall which includes rheumatoid
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Matches 756; Conservative
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The invention relates to isolated polynucleotide (I) and probes, polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polymerase chain reaction (PCR) primers, oligomers, and for chromosome and gene mapping, and in recombinant production of (II). The polynucleotides are also used in diagnostics as expressed sequence tags for identifying expressed genes. (I) is useful in gene therapy techniques to restore normal activity of (II) or to treat disease states involving contricting a polypeptide in tissue, as molecular weight markers and as a food supplement. (II) and its binding partners are useful in medical imaging of sites expressing (II). (I) and (II) are useful for treating classing of sites expressing (II). (I) and (II) are useful for treating disposptide and polynucleotide sequences have applications in diagnostics, forensics, gene mapping, identification of mutations are accounted to the polypeptical content of the polyperior of the polyperior of prettic disorders or other traits to assess products ity
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Note: The sequence data for this patent did not appear in the printed specification, but was obtained in electronic format directly from WIPO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New isolated polynucleotide and encoded polypeptides, useful in diagnostics, forensics, gene mapping, identification of mutations responsible for genetic disorders or other traits and to assess
                                                                                                                             mapping; gene mapping; gene therapy; forensic;
medical imaging; diagnostic; genetic disorder;
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Pred. No. 3.6e-191;
0; Mismatches 27; 1
                                                                                        encoding novel human diagnostic protein #22375
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Best Local Similarity 96.2'
Matches 756; Conservative
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                                                                                                                           Human; chromosome
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                                                                                                                                                                                                                                                                                                                                                                                                                            CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 717
                                                                                                                                                                                                                                                                                                        ctgaccgttaaaagacaaggactctattatatctatgcccaagtcaccttctgttccaat 579
                                                                                                                                                                                                                                                                                                                                       caaattgcggcacatgtcataagtgaggccagcagtaaaacaacatctgtgttacagtgg 459
                                                                                                                                                                                                                                                            TICGAGAGAATCITACICAGAGCIGCAAAIACCCACAGTICCGCCAAACCIIGCGGGCAA 657
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 777
                   121 CTTTTTGCTGTGTATTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180
                                  TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
                                                                                                                     340 aacaaagaggagacgaagaaagaaagactttgaaatgcaaaaaggtgatcagaatcct 399
                                                                                                                                                                                                                                                GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAAG 477
                                                                                                                                                                                                                                                                                              CIGACCGITAAAAGACAAGGACICTAITAIAITCIATGCCCAAGICACCIICTGTICCAAI 537
AACAAAGAAGA --- GAAAAAAAAAAAAAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 357
                                                                                                                                                                                                    CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 417
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human CD40L mutein coding sequence, with substitution at codon 194
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag- a
/product= CD40L_mutein
/note= "Encodes CD40L in which the Cys residue at
/note= "Encodes CD40L in which the Cys residue at
/note= position 194 is replaced by Trp, Ser, Asp
                                                                                                                                                                                                                                                                                                                                                                                              181 GAAGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAAGGAGGAGGATCTTTATCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CD40 ligand; membrane bound glyčoprotein; B cell proliferation; antibody secretion; immunoglobulin E; cytokine; mutein; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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ID AAT58122 standard; cDNA; 840 BP.
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Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         46 atgategaaacatacaaccaaactteteceegatetgeggecactggaetgeceateage 105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This sequence covers CDNA coding for human CD40L mutein polypeptides in which the wild-type Cys residue at position 194 has been replaced by a Trp. Ser. Asp or Lys residue. The muteins, designated c194W, c194S, c194D and c194K, are functional analogues of CD40L and as such are useful to induce B cell proliferation and antibody secretion. This property may be used to treat disorders associated with low levels of B cells or antibody secretion. The new CD40L muteins have a higher binding affinity for human CD40 than native CD40L and can be used in binding assays.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                    New CD40 ligand mutein with higher CD40 affinity than native ligand - useful in binding assays, and for therapy of disorders and diseases involving low levels of B cells and antibody secretion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GAAGATTTTGTÄTTCATAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 ATGATCGAAACATACAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61 ATGAAAATTTTTATGTATTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Note: The present sequence does not appear in the specification it has been produced by modifying the wild-type coding sequence, which is given on pages 19-20, at codon 194).
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                                     mutated t
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 840 BP; 266 A; 184 C; 174 G; 213 T; 3 other;
                      /*tag= b
/note= "Wild-type IGC (Cys) codon is
a codon for Trp, Ser, Asp or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 723.8; DB 18;
Pred. No. 1.8e-190;
                                                                                                                                                                                                                                                                                                                                 Spriggs MK;
                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1997-052320/05.
P-PSDB; AAW09113, AAW09114, AAW09115, AAW09116.
                                                                                                                                                                                                                                                                                                                                   Gibson MG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; Page -; 31pp; English.
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95US-0477733
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625..627
                                                                                                                                                                                                                                                                                            (IMMV ) IMMUNEX CORP.
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Best Local Similarity
Matches 753; Conserv
                                                                                                                                                                                                                                                                                                                                   Armitage RJ,
Srinivasan S;
                                                                                                         WO9640918-A2.
                                                                                                                                                                                                                                07-JUN-1995;
                                                                                                                                                                                      06-JUN-1996;
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        mutation
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          CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                       CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                                                                                                                                            TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                                 ttcgagagaatcttactcagagctgcaaatacccacagttccgccaaaccttgcgggcaa
                                                                                                                                                                                                    CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
                                        GCTGAAAAAGGATACTACACCATGAGCAACATTGGTAACCCTGGAAAATGGGAAAACAG
                                                                                                                                                                        New nucleic acid encoding human gp39 T cell antigen - which is a ligand for the CD40 receptor, causing proliferation and differentiation of B cells and some cancer cells
                                                                                                                                                                                                                                                                                                                                                                                                                                      proliferation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Aruffo A;
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22..807
/*tag= a
                                                                                                                                                                                                                                                                                                                                                                                                                                      CD40 ligand;
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                                                                                                                                                                                                                                                                                                                                                                                                                 Genomic sequence of human gp39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; Fig 1; 39pp; English
                                                                                                                                                                                                                                                                                                                                                       AAQ57984 standard; DNA; 840
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hollenbaugh D,
                                                                                                                                                                                                                                                                                                                                                                                                                                      p39; T-cell antigen; CD40 -cell differentiation; ss
                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
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P-PSDB; AAR59548.
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The complete nucleic acid sequence of human gp39 (hgp39) protein (corresp. to cDNA) and the complete AA sequence of hgp39 are presented in AAO57984 and AAR49548 respectively and contd. in plasmid CDM8-hgp39, deposited with the ATCC as E. coli, CDM8 MC1061/p3-hgp39 and assigned accession No. 69050. The human T cell antigen gp39 is a ligand for the CD40 receptor. Soluble gp39 may be produced using the
                                                                                                                                                                                                                                                                                                                                                            141
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                                                                                                                                                                                                                                                                                                                                                                                                                               142 ctttttgctgtgtatcttcatagaaggttggacaagatagaagatgaaaggaatcttcat
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                                                                                                                                                                                          Length
                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                             T; 0 other;
                                                                                                                                                                                          92.4%; Score 723.6; DB 15;
95.9%; Pred. No. 2e-190;
iive 0; Mismatches 29;
                                                                                                                                          Sequence 840 BP; 263 A; 180 C; 183 G; 214
                                                                                                       expression vector CD8-gp39.
                                                                                                                                                                                                                            Matches 754; Conservative
                                                                                                                                                                                                            Similarity
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Search completed: May 30, 2002, 03:03:47 Job time: 11276 sec ptodata/2/pna/US6007_COMB.. /ptodata/2/pna/US6008_COMB.. /ptodata/2/pna/US6009_COMB.. /ptodata/2/pna/US6010_COMB..

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Copyright (c) 1993 - 2000 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution. Description Sequence Seguence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence Sequence PCT-US94-00786-7 PCT-US94-14767-1 PCT-US97-11956-1 US-08-982-272-1 SUMMARIES DB Length Query Match Score

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658 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 717
                                                                                                                                                             GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCTTTGGCTTACTCAAA
                                                                                                 541 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
                                                                                                                                       598 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Kipps, Thomas J.
APPLICANT: Kipps, Thomas J.
APPLICANT: Sharma, Sanjai
APPLICANT: Cantwell, Mark
TITLE OF INVENTION: CONTAINING ACCESSORY
TITLE OF INVENTION: MOLECULE LIGAND GENES AND
TITLE OF INVENTION: MOLECULE LIGAND GENES AND
TITLE OF INVENTION: MODULATION AND TREATMENT OF
TITLE OF INVENTION: MALIGNANCIES
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
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MEDIUM TYPE: 3.5" Diskette,
MEDIUM TYPE: 1.44 Mb storage
COMPUTER: 18M Compatible
OPERATING SYSTEM: TBM P.C. DOS 5.0
SOFTWARE: FastSeq Version 2.0
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/08/982,272
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APPLICATION NUMBR: 60/132145
FILING DATE: 12/9/96
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1, Application US/08982272 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Guise, Jeffrey W. REGISTRATION NUMBER: 34,613
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STATE: Ca.
COUNTRY: U.S.A.
90071-2066
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California
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                                                                                  OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/107,353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 726.8; DB 5;
Pred. No. 1.3e-160;
0; Mismatches 27;
                                                                                                                                                                                        CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Particia A
RECISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 1003
TELECOMUNICATION INFORMATION:
TELECPHONE: (206)233-0644
                                                       E: Floppy disk
IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear MOLECULE TYPE: CDNA to mRNA HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 786 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 92.8°
Best Local Similarity 96.2°
Matches 756; Conservative
                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN: CD40-L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: CDS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORIGINAL SOURCE:
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                                                                            COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; LOCATION:
US-08-107-353-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ANTI-SENSE:
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TITLE OF INVENTION: DETECTION AND TREATMENT OF MUTATIONS TITLE OF INVENTION: IN A CD40 LIGAND GENE

WIDMER,

IMMUNEX CORPORATION

ADDRESSEE:

STREET:

NUMBER OF SEQUENCES: 1 CORRESPONDENCE ADDRESS:

51 UNIVERSITY STREET

STATE: WASHINGTON

USA

COUNTRY:

98101

SEATTLE

SYSTEM: Apple Operating System 7.1 MS Word for Apple 5.1, Version a PCT/US94/00786

SOFTWARE: MS WORD FOR CURRENT APPLICATION DATA:

OPERATING SYSTEM:

APPLICATION NUMBER:

OMETTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh

08/009,258

PERKINS, PATRICIA ANNE

FILING DATE: 01/22/93
ATTORNEY/AGENT INFORMATION:

PRIOR APPLICATION DATA:
APPLICATION NUMBER:

CLASSIFICATION:

FILING DATE:

REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430

LENGTH: 840 base pairs TYPE: nucleic acid . STRANDEDNESS: single

INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:

2065870606

TELEFAX:

ORGANISM: Homo sapiens

ANTI-SENSE: NO ORIGINAL SOURCE

IMMEDIATE SOURCE:

CLONE: FEATURE:

46..831

NAME/KEY:

LOCATION:

MOLECULE TYPE: CDNA HYPOTHETICAL: NO

linear

TOPOLOGY:

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TTCGAGAGAGATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGAA 660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         120
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                                                                                                                                                         61 ATGAAAATTTTTATGTATTTACTTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA
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                                                                  3;
                                       DB 13; Length 786;
                                                                    Indels.
                                         Score 726.8; DB 13;
Pred. No. 1.3e-160;
); Mismatches 27;
                                         92.8%;
96.2%;
                                                                     Matches 756; Conservative
                                                         Best Local Similarity
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CTCTGA 786
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US-08-982-272-1
                                            Query Match
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180 240 300 46 ATGATCGAAACATACAACCATCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 105 61 ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120 225 285 TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 121 CITITIGCIGIGIAICITCAIAGAAGAITGGAIAAGGICGAAGAAGGAAGIAAACCITCAI 181 GAAGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAAAGGAGAAGGATCTTTATCC 106 ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA Length 840; Indels Score 726.8; DB 1; Pred. No. 1.4e-160; 0; Mismatches 27; Ouery Match 92.8%; Best Local Similarity 96.2%; Matches 756; Conservative PCT-US94-00786-7 241 δλ g g . qa δλ qq Ω q ð ò

Sequence 7, Application PC/TUS9400786 GENERAL INFORMATION:

RESULT 4 PCT-US94-00786-7

RICHARD VILLIAM

ARMITAGE, FANSLOW, RENSHAW, DAVISON,

APPLICANT: APPLICANT: APPLICANT: APPLICANT: APPLICANT

BLAIR MELANIE

SPRIGGS,

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                                                                                                                        CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT 537
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301 AACAAAGAAGA---GAAAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 357
                                                                                                           GCTGAAAAAGGATACTACACCATGAGCAACATTGGTAACCCTGGAAAATGGGAAACAG 477
                  TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: LONGO, DAN L.
APPLICANT: MURRHY, WILLIAM
TITLE OF INVENTION: METHOD OF PREVENTING OR TREATING
TITLE OF INVENTION: DISEASE CHARACTERIZED BY NEOPLASTIC CELLS
TITLE OF INVENTION: EXPRESSING CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Microsoft Word for Macintosh, Version #5.1a CURRENT APPLICATION DATA:
APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USSN 08/172,664
er 23, 1993
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 1, Application PC/TUS9414767 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IMMUNEX CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           December 21, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: PERKINS, PARLICIA A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2818
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: LONGO, DAN
APPLICANT: MURPHY, WILLIAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: December 23,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA: APPLICATION NUMBER: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES:
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WASHINGTON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
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                                                                                                                                                                                                                                                                                                                                                                                     Length 840;
                                                                                                                                                                                                                                                                                                                                                                                    Score 726.8; DB 1;
Pred. No. 1.4e-160;
0; Mismatches 27;
TELECOMMUNICATION INFORMATION:
            TELEPHONE: (206)587-0430
TELEFAX: (206)333-0644
INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                    92.8%;
ilarity 96.2%;
Conservative
                                                                                 LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                         ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                                                                                                                                                                                                                                 46..831
                                                                                                                                        linear
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S
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ORIGINAL SOURCE:
                                                                                                                    STRANDEDNESS:
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HYPOTHETICAL: 1
                                                                                                                                        TOPOLOGY:
                                                                                                                                                                                                                                                                                               ; NAME/KEY:
; LOCATION:
PCT-US94-14767-1
                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Simi
Matches '756;
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FEATURE:
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GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG 477
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 658 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 717
                                                                                                                                                                                    241 TIGCIGAACIGIGAGGAGAIGAGAAAGGCAATITGAAGACCIIGICAAGGAIAIAACGITA 300
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                                                                                                                                                                                                                                                                                                                                                                                                                                              61 ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA
                478 CIGACCGITAAAAGACAAGGACTCTATTATATTTTATGCCCAAGTCACCTTCTGTTCCAAT
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                                                             CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT
                                                                                                                       301 AACAAAGAAGA---GAAAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                                                                                                                                                                             CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAAACAACATCTGTGTTACAGTGG
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                    Length 840;
                                                                                                                                                                                    Sequence 1, Application PC/TUS9711956
GENERAL INFORMATION:
APPLICANT: Immunex Corporation
APPLICANT: Immunex METHOD OF ACTIVATING DENDRITIC CELLS
NUMBER OF SEQUENCES: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                       Version #6.0.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 726.8; DB 1;
Pred. No. 1.4e-160;
0; Mismatches 27;
                                                                                                                                                                                                                                                                                                                                                                                                                         SYSTEM: Apple Operating System Microsoft Word for Macintosh, V
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/677,762
FILING DATE: 10-JUL-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/763,995
FILING DATE: 12-DEC-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCI/US97/11956
FILING DATE: 09-JUL-1997
                                                                                                                                                                                                                                                                                                                                                             ZIP: 98101
COMPUTER READABLE FORM:
WEDLUM TYPE: Floppy disk
COMPUTER: Power Macintosh 7200/90
OPERATING SYSTEM: Apple Operating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Perkins, Patricia A. REGISTRATION WUMBER: 34,693 REFERENCE/DOCKET NUMBER: 2845-WO TELECOMMUNICATION INFORMATION: TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                 NOTRESSEE: IMMUNEX CORPORATION STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92.88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (206)233-0644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
                                                                                                                                                                                                                                                                                                                                  WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                   SEATTLE
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                                                                                   778 CTCTGA 783
                                                                                                                826 CTCTGA 831
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ANTI-SENSE: N
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PCT-US97-11956-1
                                                                                                                                                             RESULT 6
PCT-US97-11956-1
                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
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NOVEL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                  5: IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                             Sequence 10, Application US/07805723 GENERAL INFORMATION:
                                                                                                                               APPLICANT: ARMITAGE, RICHARD APPLICANT: FANSLOW, WILLIAM APPLICANT: SPRIGGS, MELANIE
                                                                                                                                                                                TITLE OF INVENTION: NO NUMBER OF SEQUENCES: 1 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                  CITY: SEATTLE
STATE: WASHINGTON
COUNTRY: 112A
98101
                                                                                                                                                                                                                                         ADDRESSEE:
                                                                 RESULT 7
US-07-805-723-10
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1 ATGATGGAAACATACAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60

96.2%;

Similarity

Conservative

756;

Best Local Matches 75

Gaps

Indels

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Gaps
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                                                                                                                            718 GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA
                                                     658 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PAPPLICATION NUMBER: US/07/969,703A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 726.8; DB 3;
Pred. No. 1.4e-160;
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                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
TITLE OF INVENTION: NOVEL CYTOKINE
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                               US-07-969-703A-11; Sequence 11, Application US/07969703A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                :: IMMUNEX CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              34,693
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TELEPHONE: 2065870430
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INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARATTERISTICS:
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NAME: Perkins, Patricia A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 34, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            : 840 base pairs
NUCLEIC ACID
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MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: SEATTLE
STATE: WASHINGTON
COUNTRY: USA
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Best Local Similarity
Matches 756; Conservat
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HYPOTHETICAL: N
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ORIGINAL SOURCE:
ORGANISM: HOM
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US-07-969-703A-11
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                                                                                                                                                                                                                                                                                              RESULT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 645
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           301 AACAAAGAAGA---GAAAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT 357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 417
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   526 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT 585
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 840;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 92.8%; Score 726.8; DB 3; Best Local Similarity 96.2%; Pred. No. 1.4e-160; Matches 756; Conservative 0; Mismatches 27;
PatentIn Release #1.0, Version #1.25
                                APPLICATION NUMBER: US/07/805,723
                                                                                                                                       REFERENCE/DOCKET NUMBER: 2E
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                                                                                            32585
                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS:
                                                                                ATTORNEY AGENT INFORMATION:
NAME: OSTER, JEFFREY B.
REGISTRATION NUMBER: 3256
                                                                                                                                                                                                                                                        840 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
CLONE: CD40-L
               CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                              single
                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: CDNA
                                                                                                                                                                                                                                                                                                               linear
                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                   ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                          STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                              TOPOLOGY:
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                                                                                                                                                                                                                                                          LENGTH:
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                                                                                                                                                                    GAAGATTTTGTATTCATAAAAAAGCTAAAGATGCAACAAAGGAGGAGGATGTTTATCC 240
                                                                                                                                                                                                                                    241 TIGCIGAACIGIGAGGAGAIGAGAGGCAATITGAAGACCITGICAAGGAIAIAAACGITA 300
                                                                                                                                                                                                                                                                 286 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 345
                                                                                                                                                                                                                                                                                                                                                                  CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 417
                                                                                                                                                                                                                                                                                                                                                                                                                                 418 GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG 477
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      718 GTGACTGATCCAAGCCAAGTGAGCCATGGCACTGGCTTCACGTCCTTTGGCTTACTCAAA 777
46 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 105
                                                                                                                                                                                        GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG
                               61 ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA
                                                                                                                                                                                                                                                                                                     .AACAAAGAAGA - - -GAAAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
TITLE OF INVENTION: NOVEL CYTOKINE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 11, Application US/07969703B GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IMMUNEX CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 17 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WASHINGTON
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                         Patentin Release #1.0, Version #1.25
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Pred. No. 1.4e-160;
); Mismatches 27;
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                                                           US/07/969,703E
              PC-DOS/MS-DOS
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IBM PC compatible
                                                                                       CLASSIFICATION: 435
ATTORNEY/AGENET INFORMATION:
NAME: PERKINS, PALTICIA A.
REGISTRATION NUMBER: 34,693
                                                                                                                                                                  TELECOMMUNICATION INFORMATION: TELEPHONE: 2065870430
TELEFAX: 2065870606
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llarity 96.2%;
Conservative
                                                                                                                                                     REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens
                             SOFTWARE: Patentin Rel CURRENT APPLICATION DATA:
                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                          APPLICATION NUMBER: US
FILING DATE: 19921023
                                                                                                                                                                                                                                                                                          linear
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hes 756; Conserv
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ORGANISM: HOM
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HYPOTHETICAL:
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                                                     TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                                                                                                                                                                                             APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: WIDDER, MELANIE
APPLICANT: WIDDER, MICHAEL
TITLE OF INVENTION: DETECTION AND TREATMENT OF MUTATIONS
TITLE OF INVENTION: IN A CD40 LIGAND GENE
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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FILING DATE: 19930122
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STREET: 51 UNIVERSITY STREET
CITY: SEATTLE
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US-08-009-258-7
Sequence 7, Application US/08009258
; GENERAL INFORMATION:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
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ATTORNEY/AGENT INFORMATION:
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TYPE: NUCLEIC ACID
STRANDEDNESS: single
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IMMEDIATE SOURCE:
CLONE: CD40-L
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TELEFAX: 2065870606
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MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
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CLASSIFICATION:
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US-08-009-258-7
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                                           Gaps
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TITLE OF INVENTION: DISEASE CHARACTERIZED BY NEOPLASTIC CELLS
TITLE OF INVENTION: EXPRESSING CD40
    Length
                                         Indels
                                         27;
Score 726.8; DB 4;
Pred. No. 1.4e-160;
                                         0; Mismatches
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GENERAL INFORMATION:
APPLICANT: ARMITAGE, RICHARD
APPLICANT: FANSLOW, WILLIAM
APPLICANT: LONGO, DAN L.
APPLICANT: MURPHY, WILLIAM
92.8%;
96.2%;
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MURPHY, WILLIAM
                                         Matches 756; Conservative
                     Similarity
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US-08-172-664-1
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406 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 465
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                                418 GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAACAG
                                                                                                  478 CIGACCGITAAAAGACAAGGACICIAİTAIAICIAIGCCCAAGICACCIICIGIICCAAI
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TITLE OF INVENTION: Method of Diagnosing or Treating Hodgkin'
TITLE OF INVENTION: Disease
NUMBER OF SEQUENCES: 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SYSTEM: Apple Macintosh System 7.1
Microsoft Word for Macintosh, Version #5.1a
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APPLICATION NUMBER: US/08/215,862
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSE: IMMUNEX CORPORATION
STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Perkins, Parricia A. REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 2820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 1, Application US/08215862 GENERAL INFORMATION: APPLICANT: Gruss, Hans-J_rqen
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Floppy disk COMPUTER: Macintosh'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
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nucleic acid
DEDNESS: single
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COMPUTER READABLE FORM:
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STATE: WASHINGTON
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HYPOTHFFTTT
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CITY: SE
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                                                                                                                                                                                            OPERATING SYSTEM: Apple Macintosh System 7.1
SOFTWARE: Microsoft Word for Macintosh, Version #5.1a
CURRENT APPLICATION DATE: US/08/172,664
FILING DATE: December 23, 1993
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Pred. No. 1.4e-160;
0; Mismatches 27;
         NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS: :
STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                     NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2818
                                                                                                                                                                                                                                                                                                                                                                        TELECOMMUNICATION INFORMATION: (206)587-0430
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Best Local Similarity 96.2%;
Matches 756; Conservative
                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
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IMMEDIATE SOURCE:
CLONE: CD40-L
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nucleic acid
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US-08-172-664-1
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US-08-234-580-3
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                                                                        Length 840;
                                                                        92.8%; Score 726.8; DB 6;
llarity 96.2%; Pred. No. 1.4e-160;
Conservative 0; Mismatches 27;
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                                     46..831
                                                                        Query Match
Best Local Similarity
Matches 756; Conserv
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IMMEDIATE SOURCE
        CD40-L
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US-08-215-862-1
         CLONE:
                 FEATURE:
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TITLE OF INVENTION: METHODS FOR PROLIFERATING AND TITLE OF INVENTION: DIFFERENTIATING B CELLS, AND UNMBER OF SEQUENCES: 4
NUMBER OF SEQUENCES: ADDRESS:
ADDRESSER
                                                                                                                                                                                                                                                                   Version #1.25
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Pred. No. 1.4e-160;
0; Mismatches 27;
                                                                                                 ADDRESSEE: STERN, KESSLER, GOLDSTEIN & FOX STREET: 100 NEW YORK AVE. N.W. SUITE 600 CITX: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 1011.1030000/RAM TELECOMMUNICATION INFORMATION: ... TELEPHONE: (202)371-2600
                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/234,580
FILING DATE: 28-APR-1994
CLASSIFICATION: 435
                                                                                                                                                                                                            MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                           36,217
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96.2%;
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NAME: MILLMAN, ROBERT A
REGISTRATION NUMBER: 36,
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LENGTH: 840 base pairs
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STATE: D.C.
TO 20005
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                                                                502 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                 598 TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: Apple Power Macintosh System 7.5.5 SOFTWARE: Microsoft Word for Macintosh, Version 6.0.1 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: CAMPBELL, KIM A.
APPLICANT: KENNEDY, MARY K.
APPLICANT: KENNEDY, MARY K.
APPLICANT: MALISZEWKII, CHARLES R.
TITLE OF INVENTION: METHOD FOR STINULATING I
TITLE OF INVENTION: PROTEIN
FILLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        IMMUNEX CORPORATION
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application US/08396230 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      34,693
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TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: March 1, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TYPE: nucleic acid
STRANDEDNESS: single
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ORIGINAL SOURCE:
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US-08-396-230-1
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                                                                                                                                                                   46 ATGATCGAAACATACAACCATCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 105
                                                                                                                  Gaps
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                                                                                                                                              1 ATGATCGAAACATACAACCAAACTTCTCCCCGGATCTGCGGCCACTGGACTGCCCATCAGC
                                                                                Length 840;
                                                                                                                  Indels
                                                                     Score 726.8; DB 7;
Pred. No. 1.4e-160;
                                                                                                                    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 11, Application US/08477733A GENERAL INFORMATION:
                                                                                   .92.8%;
96.2%;
                                                                                                                      Conservative
                  46..831
                                                                                                  Similarity
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826 CTCTGA 831
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, NAME/KEY:
; LOCATION:
US-08-396-230-1
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APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIYASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
CORRESPONDENCES: 24
CORRESSEE: IMMINIER ADDRESSE:
                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, version 5.1a
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/08/477,733A FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: 08/249,189
FILING DATE: May 24, 1994
CLASSIFICATION: 435
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APPLICATION NUMBER: 07/783,707
FILING DATE: October 25, 1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: PERKIALS, PATITOLS A.
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
TELEPHONE: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    October 23, 1992
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MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 840 base pairs
TYPE: nucleic acid
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IMMEDIATE SOURCE:
CLONE: CD40-L
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PRIOR APPLICATION DATA:
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APPLICATION NUMBER: (
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CITY: SEATTLE
STATE: WASHINGTON
COUNTRY: USA
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LOCATION: 46..831
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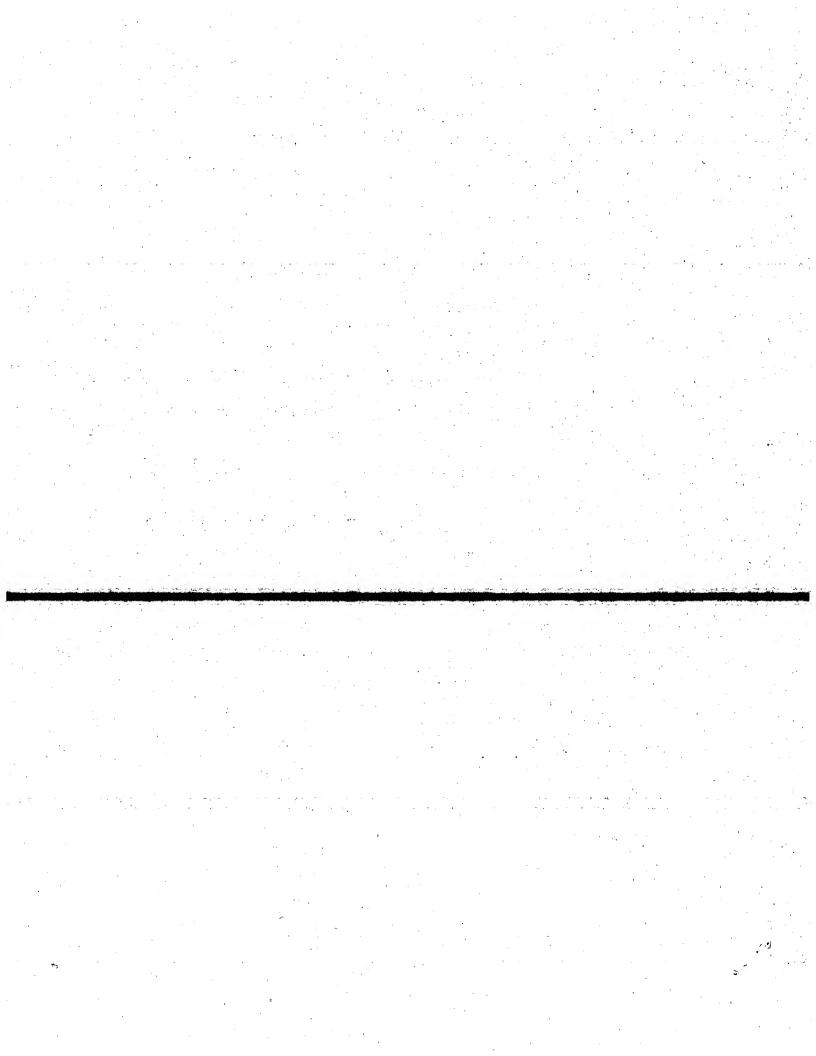
Indels

Best Local Similarity 96.2 Matches 756; Conservative

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Query Match

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CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180
                                                                                              GAAGATTTTGTATTCATAAAAAAGCTAAAGAGATGCAACAAAGGAGAAGGATCTTTATCC 240
                                                                                                                                                                          241 TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
                                                                                                                                                                                                                                                                                                                                                                         358 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 417
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326 CTCTGA 831
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Search completed: May 30, 2002, 02:52:00 Job time: 11624 sec

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597
                                          537
                                                         538 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCGGTAGA
                                        CTGACCGTTAAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OPERATING SYSTEM: Apple Operating System 7.1 SOFTWARE: Microsoft Word for Apple, version 5.1a CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/770,974
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
NUMBER OF SEQUENCES: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-DEC-1996
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FILING DATE: October 25, 1991
ATTORNEY/AGENT INFORMATION:
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IOR APPLICATION PART
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,733
FILING DATE: 02-AUG-1995
APPLICATION NUMBER: 08/249,189
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December 5, 1991
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                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 11, Application US/08770974
Patent No. 6290972
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                              ARMITAGE, RICHARD
FANSLOW, WILLIAM
SPRIGGS, MELANIE
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MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh.
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APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                             ADDRESSEE
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US-08-770-974-11
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TTCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 657
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                                                                                                                                                                                                                                                                                                                                                                                           Length 840;
                                                                                                                                                                                                                                                                       Score 726.8; DB 4;
Pred. No. 1e-197;
0; Mismatches 27;
                                                                                                                                                                                                                                                                           92.8%;
             INFORMATION FOR SEQ 1D NO: 1 SEQUENCE CHARACTERISTICS: LENGTH: 840 base pairs TYPE: nucleic acid
                                                                                                                                                            ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                      Matches 756; Conservative
2065870430
2065870606
                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                             TOPOLUGITYPE: CD
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      TELEPHONE:
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US-08-770-974-11
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                                                                           301 AACAAAGAAGA---GAAAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                                                                                                                           558 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
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APPLICANT: SPRIGGS, MELANKE
APPLICANT: STRINYASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TILLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.1
SOFTWARE: Microsoft Word for Apple, version 5.1a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/769,819
FILING DATE: 19-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 11, Application US/08769819 Patent No. 6264951
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: ARMITAGE, RICHARD APPLICANT: FANSLOW, WILLIAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
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                   08/249,189
                                                                           07/969,703
                                                                                                                                   07/805,723
                                                                                                                                                                                         07/783,707
                                                                                         FILING DATE: October 23, 1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
                                                                                                                                                                                                         FILING DATE: October 25, 1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                             NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 28
                                                                                                                                                                                                                                                                                                                                                              11:
                                   FILING DATE: May 24, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/965
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92.88;
07-JUN-1995
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TYPE: nucleic acid
STRANDEDNESS: single
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Matches 756; Conservative
                                                                                                                                                                                                                                                                                                                  TELEPHONE: 2065870430
TELEFAX: 2065870606
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                 APPLICATION NUMBER:
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; LOCATION:
US-08-769-819-11
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418 GCTGAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG 477

358 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGŤGG

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                                                                                             Score 726.8; DB 3;
Pred. No. 1e-197;
0; Mismatches 27;
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US-08-589-771B-7
; Sequence 7, Application US/08589771B
                                                                                                tch 92.8%;
al Similarity 96.2%;
756; Conservative
                                                    46..831
                                                                                                Query Match
Best Local Similarity
Matches 756; Conserv
ORGANISM: HOMO IMMEDIATE SOURCE: CLONE: CD40-L
                                        , NAME/KEY: CDS
, LOCATION: 46.
US_09-088-913A-11
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CTCTGA 831
                                 FEATURE
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                                                                                                 TREATMENT OF INDIVIDUALS EXHIBITING
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Pred. No. 1e-197;
0; Mismatches 27;
                                                                                                                    DEFECTIVE CD40L (as amended)
                                                                                                                                                                                                                                                                                                                                        UMBER: US/08/589,771B
January 22, 1996
                                                                                                                                                                                                                                                                                                     SYSTEM: MS-DOS/Window Word for Windows 95,
                                                                                                                                                             ADDRESSEE: IMMUNEX CORPORATION STREET: 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                           08/009,258
                                                                                                                                                                                                                                                                       E: Floppy disk
IBM PC Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    34,347
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                            BARRY
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96.2%;
                                                                         MELANIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: Nucleic acid STRANDEDNESS: single TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY AGENT INFORMATION:
NAME: HENRY, JANIS C.
REGISTRATION NUMBER: 34,3
                                                       BLAIR
                                                                                      WIDMER, MICHAEL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 96.2
Matches 756; Conservative
                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA: APPLICATION NUMBER: US,
                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08
FILING DATE: 01/22/93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2065870430
                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                FILING DATE: Januar CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2065870606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           46..831
                                                                       SPRIGGS,
                                                                                                                                                                                                             STATE: WASHINGTON COUNTRY: USA
                                                                                                                                     NUMBER OF SEQUENCES:
                                           FANSLOW,
                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
                                                                                                    TITLE OF INVENTION:
                                                         RENSHAW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo
                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IMMEDIATE SOURCE:
GENERAL INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORIGINAL SOURCE:
                                                                                                                                                                                               CITY: SEATILE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HYPOTHETICAL:
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                                                                                                                                                                                                                                               98101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
                                                                                                                                                                                                                                                                                          COMPUTER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ANTI-SENSE:
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                                                                         APPLICANT:
                                                                                         APPLICANT:
                                                           APPLICANT:
               APPLICANT
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                              APPLICANT
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706 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTGGGTGTTTGTCAAT 765
                                                                                     718 GTGACTGATCCAAGCCAAGTGAGCCATGGCATTGACGTCTTTGGCTTACTCAAA 777
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 98101
COMPUTER READABLE FORM:
MEDIUM TYEE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: MS Word for Apple Power Macintosh, version 6.0.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: SRINIVASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
APPLICANT: MORRIS, ARVIA E.
APPLICANT: MCGREW, JEFERRY
ITLE OF INVENTION: NOVEL CYTOKINE THAT BINDS CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/088,913A
                                                                                                                                                                                                                                                                               Sequence 11, Application US/09088913A Patent No. 6087329 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
APPLICATION NUMBER: 08/477,733
FILING DATE: June 07, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/969,703
FILING DATE: October 23, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: IMMUNEX CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/783,707
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/484,624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: October 25, 1991 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION
TELEPHONE: 2065870430
TELEFAX: 2065870606
                                                                                                                                                                                                                                                                                                                                                APPLICANT: ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                FANSLOW, WILLIAM SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: NO NUMBER OF SEQUENCES: 2 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 51 UNIVERS
CITY: SEATTLE
STATE: WASHINGTON
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                                                                                                                                             778 CTCTGA 783
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                               US-09-088-913A-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   357
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    538 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA 597
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  658 CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 717
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              358 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 840;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 726.8; DB 3; Length Pred. No. 1e-197; O; Mismatches 27; Indels
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2845
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206)587-0430
TELEPHONE: (206)587-0430
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    92.8%;
96.2%;
                                                                                                                                                                                                                                                           ANTI-SENSE: NO
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 96.2<sup>3</sup>
Matches 756; Conservative
                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: cDNA
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                      46..831
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                                                                                                                                                                                                                                                                                                                                                                                   NAME/KEY: CDS
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US-08-763-995-1
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226 GAAGAITITGIAITCATGAAAACGATACAGAGATGCAACACAGGAGAAAGAICCTTAICC 285
                                TTGCTGAACTGTGAGGAGTGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300
                                                     286 TTACTGAACTGTGAGGAGATTAAAAGCCAGTTTGAAGGCTTTGTGAAGGATATAATGTTA 345
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                                                                                                                      CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG
                                                                                                                                                                                      418 GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG
                                                                                                                                                                                                                                                            466 GCTGAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAAAACAG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: MARASKOVSKY, EUGENE
ATTILE OF INVENTION: METHOD OF ACTIVATING DENDRITIC CELLS
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Apple Operating System
ft Word for Macintosh,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UMBER: USSN 08/677,762
10 JUL 1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: Power Macintosh 7200/90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              us/08/763,995
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51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/08763995
Patent No. 6017527
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Microsoft WC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
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CITY: SEATTLE.
STATE: WASHING
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826 CTCTGA 831.
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                                                                                                                                      SYSTEM: Apple Operating System 7.5.5
MS Word for Apple Power Macintosh, version 6.0.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 726.8; DB 2;
Pred. No. 1e-197;
); Mismatches 27;
                                                                                                                                                                                              US/08/477,733B
                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
                                                                                                                                                                                                                                                                08/249,189
                                                                                                                                                                                                                                                                                                                                           FILING DATE: October 23, 1992 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: 07/783,70 FILLING DATE: October 25, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Perkins, Patricia A.
MATION NUMBER: 34,693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 28
TELECOMMUNICATION INFORMATION: 2065870430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       92.8%;
                                                                                                                                                                                                            June 07, 1995
                                                                                                                        COMPUTER: Apple Macintosh OPERATING SYSTEM: Apple O
                                                                                                                                                                                                                                                                              May 24, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                            URRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     nucleic acid
EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION DATA:
                                                                                                                                                                                                                              CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2065870606
                                                                                         COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER:
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                                         WASHINGTON
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Best Local Similarity
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CLONE: CD40-L
                                                                                                                                                                                                                                                                                                FICATION:
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                                                           USA
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OPERATING SYSTEM: Apple Operating System 7.5.5
SOFTWARE: MS Word for Apple Power Macintosh, version 6.0.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,624A
                                                                                                                 NOVEL CYTOKINE THAT BINDS CD40
                                  SRINIVASAN, SUBHASHINI
GIBSON, MARYLOU
                                                                                                                                                                             : IMMUNEX CORPORATION
51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/477,733
FILING DATE: June 07, 1995
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APPLICATION NUMBER: 07/805,723
FILING DATE: December 5, 1991
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APPLICATION NUMBER: 07/783,707
FILING DATE: OCtober 25, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: October 23, 1992 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia A.
REGISTRATION NUMBER: 34,693
REFERENCE/COCKET NUMBER: 2802
TELECOMMUNICATION INFORMATION:
TELEPHONE: 2065870430
FANSLOW, WILLIAM
SPRIGGS, MELANIE
                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
                                                                          APPLICANT: MORRIS, ARVIA E. APPLICANT: 'MCGREW, JEFFERY TITLE OF INVENTION: NOVEL C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 840 base pairs
TYPE: nucleic acid
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                                                                                                                                    NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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ORIGINAL SOURCE:
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HYPOTHETICAL: N
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US-08-484-624A-11
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APPLICANT:
APPLICANT:
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GENERAL INFORMATION:
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GIBSON, MARYLOU
MORRIS, ARVIA E.
MCGREW, JEFFERY
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CORRESPONDENCE ADDRESS:
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APPLICANT:
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0; Mismatches Score 726.8; Pred. No. 1e-

92.8%; nilarity 96.2%; Conservative

Ouery Match Best Local Similarity Matches 756; Conserva

Length 840;

8; DB 2; Length 1e-197; thes 27; Indels

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IMMEDIATE SOURCE:
CLONE: CD40-L
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                                         , NAME/KEY:
; LOCATION:
US-08-249-189-11
                               FEATURE
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   622 ITCGAGAGAATCTTACTCAGAGCTGCAAATACCCACAGTTCCGCCAAACCTTGCGGGCAA 681
                                  CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT
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SOFTWARE: Microsoft Word for Apple, version 5.1a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/249,189
                                                                                                                                                                                                                                                                                                                                        APPLICANT: FANSLOW, WILLIAM
APPLICANT: SPRIGGS, MELANIE
APPLICANT: SRINIYASAN, SUBHASHINI
APPLICANT: GIBSON, MARYLOU
TITLE OF INVENTION: NOVEL CYTOKINE
NUMBER OF SEQUENCES: 23
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                                                                                                                                                                                                                                                                         sequence 11, Application US/08249189
Patent No. 5961974
GENERAL INFORMATION:
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APPLICATION NUMBER: 07/969,70:
FILING DATE: October 23, 1992
CLASSIFICATION: 514
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APPLICATION NUMBER: 07/783,70
FILING DATE: October 25, 1991
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FILING DATE: December 5, 1991
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                                                                                                                                                                                                                                                                                                                            ARMITAGE, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TTORNEY/AGENT INFORMATION:
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TYPE: nucleic acid
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                                                                                                                                                                  778 CTCTGA 783
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802 CTCTGA 807
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                                                                                                                                                                    ATGAAAATTTTATGTATTTACTTACTGTTTTTTTTTATCACCCAGATGATTGGGTCAGCA 120
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                                                                                                                1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC 60
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                                                          Length 840;
                                                                                        Indels
                                                           Score 726.8; DB 2;
Pred. No. 1e-197;
0; Mismatches 27;
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; sequence 11, Application US/08484624A
; Patent No. 5962406
                                                                92.8%;
ilarity 96.2%;
Conservative
CDS
46..831
                                                                Query Match
Best Local Similarity
Matches 756; Conserv
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CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAAGAAGTAAACCTTCAT 180
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                                                                                                                                                                                     CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 417
                                                                                                                                                                                                                                  GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG: 477
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                                                        AACAAAGAAGA - - - GAAAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                                                               GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG
                                                                                                                                                                                                                                                                                CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
                                                                                                                                                                                                                                                                                                                                CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
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SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CD40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/08690096
Patent No. 5945513
GENERAL INFORMATION:
APPLICANT: ARUFFO, ALEJANDRO
APPLICANT: HOLLENBAUGH, DIANE
APPLICANT: LEDBETTER, JEFFREY A.
TITLE OF INVENTION: SOLUBLE LIGANDS FOR CDA
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
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Pred. No. 1e-197;
0; Mismatches 27;
                                                                                   US 07/940,605
                                                                                                                                          REGISTRATION NUMBER: 18,872
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 5524-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212 790-9090
TELERX: 212 869-864/9741
TELEX: 6141 PENNIE
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYRE: nucleic acid
STRANDENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92.8%;
                                                                                                   FILING DATE: 04-SEP-1992
ATTORNEY/AGENT INFORMATION:
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Best Local Similarity 96.28
Matches 756; Conservative
                                                         PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                       CLASSIFICATION: 536
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22..807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; NAME/KEY:
; LOCATION:
US-08-690-096-1
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61 AIGAAAATITTAAGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
                                                          CAATCCATTCACTTGGGAGGAGTATTTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 717
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Pred. No. 1e-197;
0; Mismatches 27; Indels 3
                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: METHODS FOR PROLIFERATING AND TITLE OF INVENTION: DIFFERENTIATING B CELLS, AND USES THEREOF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                       3: STERN, KESSLER, GOLDSTEIN & FOX 100 NEW YORK AVE. N.W. SUITE 600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/431,055
FILING DATE: 28-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                              Sequence 3, Application US/08431055
Sequence 3, Application US/08431055
Patent No. 5817516
GENERAL INFORMATION HEAPLIAN R
APPLICANT: CASTLE, BRIAN E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           36,217
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TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: D.C.
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (202) 371-2540 INFORMATION FOR SEQ ID NO:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: STERN, KE
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                                                                                                                                   778 CTCTGA 783
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                                                                                                                                                                                                                                                                                                                                                                                                              STREET:
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Pred. No. 1e-197;
0; Mismatches 27; Indels 3.
                                                  2818-A
                  NAME: Perkins, Patricia A. REGISTRATION NUMBER: 34,693 REFERENCE/DOCKET NUMBER: 28 TELECOMMUNICATION INFORMATION: TELEPHONE: (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                     92.8%;
   ATTORNEY/AGENT INFORMATION:
                                                                                                             INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 840 base pairs
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 96.2
Matches 756; Conservative
                                                                                               (206)233-0644
                                                                                                                                                                                STRANDEDNESS: single
                                                                                                                                                                                                              MOLECULE TYPE: CDNA
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                                                                                                                                                                                              linear
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IMMEDIATE SOURCE:
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NO
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                                                                                                                                                                                                                                                                                                                                                            LOCATION:
US-08-360-923A-1
                                                                                                                                                                                                    POPOLOGY:
                                                                                                   TELEFAX:
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358 CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG 417
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                                                                                                                                                             418 GCTGAAAAAGGATACTACACCATGAGCAACAACTTGGTAACCCTGGAAAATGGGAAACAG
                                                                                                                                                                                                                                                                                                 CTGACCGTTAAAAGACAAGGACTCTATTATATCTATGCCCAAGTCACCTTCTGTTCCAAT
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                                                                                  718 GIGACIGAICCAAGCCAAGIGAGCCAIGGCACIGGCIICACGICCIIIGGCIIACICAAA
                                                            301 AACAAAGAAGA---GAAAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
                                                                                                                                                                                                                                                                                                                                                                       538 CGGGAAGCTTCGAGTCAAGCTCCATTTATAGCCAGCCTCTGCCTAAAGTCCCCCGGTAGA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM: Apple Macintosh System 7.1
SOFTWARE: Microsoft Word for Macintosh, Version #5.1a
CURRENT APPLICATION DATA:
FAPLICATION NUMBER: US/08/360,923A
FILING DATE: December 21, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            METHOD OF PREVENTING OR TREATING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: USSN 08/172,664
FILING DATE: December 23, 1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/08360923A Patent No. 5674492
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         E: IMMUNEX CORPORATION 51 UNIVERSITY STREET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FANSLOW, WILLIAM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: MURPHY, WILLIAM
TITLE OF INVENTION: METHOD
TITLE OF INVENTION: DISEASE
TITLE OF INVENTION: EXPRESS
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CORRESPONDENCE ADDRESS:
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COMPUTER READABLE FORM:
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CITY: SE
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                                  APPLICANT: SPRIGGS, MELANIE
APPLICANT: WIDMER, MYCHAEL
TITLE OF INVENTION: DETECTION AND TREATMENT OF MUTATIONS
TITLE OF INVENTION: IN A CD40 LIGAND GENE
NUMBER OF SEQUENCES: 16
                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: Apple Operating System 7.1 SOFTWARE: MS WORD for Apple 5.1, Version a CURENT APPLICATION DATA: APPLICATION NUMBER: US/08/184,422
                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/009,258
FILING DATE: 01/22/93
ATTORNEY/AGENT INFORMATION:
NAME: PERKINS, PATRICIA ANNE
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 2810-A
TELEPHONE: 2065870430
TELEPAX: 2065870606
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                       E: IMMUNEX CORPORATION 51 UNIVERSITY STREET
                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Apple Macintosh
OPERATING SYSTEM: Apple O
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TYPE: nucleic acid
STRANDEDNESS: single
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IMMEDIATE SOURCE:
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ADDRESSEE: IMMUNEX C
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FANSLOW,
RENSHAW,
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; LOCATION: 46..831
US-08-184-422-7
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COUNTRY:
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SYSTEM: PC-DOS/MS-DOS
PatentIn Release #1.0, Version #1.25
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STREET: 1155 Avenue of the Americas
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APPLICANT: HOLLENBAUGH, DIANE
APPLICANT: HOLLENBAUGH, DIANE
APPLICANT: LEDBETTERE, JEFFREY A.
TITLE OF INVENTION: SOLUBLE LIGANDS INUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/07/940,605A FILING DATE: 04-SEP-1992
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ALAME: MISTOCK, S. Leslie
REGISTRATION NUMBER: 18,872
REFERENCE/DOCKET NUMBER: 5624
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212 790-9090
TELERAX: 212 869-8864/9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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nucleic acid
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MOLECULE TYPE:
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781 CTCTGA 786
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658 CAATCCATTCACTTGGGAGGAGTATTGAATTGCAACCAGGTGCTTCGGTGTTTGTCAAT 717
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                                                                                                                61 AIGAAAATTTTAATGTATTTACTTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA
                                                                                                                              121 CTTTTTGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT
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Length 840;
                                Indels
    Score 726.8; DB Pred. No. 1e-197, 0; Mismatches
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; Sequence 7, Application US/08184422
; Patent No. 5565321
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DAVISON, BARRY
       Query Match 92.8%;
Best Local Similarity 96.2%;
Matches 756; Conservative
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APPLICANT: ARMITAN
APPLICANT: DAVISOI
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301 AACAAAGAAGA---GAAAAAAGAAAACAGCTTTGAAATGCAAAAAGGTGATCAGAATCCT
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                                                                                       Gaps
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                                                                                                                1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGCACTGCCCATCAGC
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GENERAL INFORMATION:
APPLICANT: Springs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
                                                             Length 786;
                                                                                        Indels
                                                            Score 726.8; DB 1;
Pred. No. 9.9e-198;
0; Mismatches 27;
                                                             92.8%;
96.2%;
                                                                                       Conservative
                                                            Query Match
Best Local Similarity
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781 CTCTGA 786
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; NAME/KEY:
; LOCATION:
US-08-446-922-3
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                                                                                     Matches
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61 ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
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                                                                                                                                                                                                                                                OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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Pred. No. 9.9e-198;
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                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: PCT/US93/10034 FILING DATE:
                                                        ADDRESSEE: Immunex Corporation
STREET: 51 University Street
CITY: Seattle
                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY AGENT INFORMATION:
NAME: Perkins, Patricia A
REGISTRATION UNDRER: 34,693
REERENCE/DOCKET NUMBER: 1003
TELECOMMUNICATION INFORMATION:
TELEFONE: (206)293-0644
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92.8%;
96.2%;
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TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity 96.2
Matches 756; Conservative
               NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Immunex C
                                                                                                                                                                                     COMPUTER READABLE FORM:
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INVENTION:
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LOCATION: 1...
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                                                                                                                                             USA
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MOLECULE TYPE:
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                                                                                                                                                                 ZIP: 98101
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                                                                                                                     STATE: W
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CAAATTGCGGCACATGTCATAAGTGAGGCCAGCAGTAAAACAACATCTGTGTTACAGTGG

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Seattle
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                                                                                                           May 30, 2002, 02:51:53; Search time 139.75 Seconds (without alignments) 1376.251 Million cell updates/sec
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... /cgn2_6/ptodata/2/ina/6B_COMB.seq:*
... /cgn2_6/ptodata/2/ina/6A_COMB.seq:*
... /cgn2_6/ptodata/2/ina/6B_COMB.seq:*
... /cgn2_6/ptodata/2/ina/PeTUS_COMB.seq:*
... /cgn2_6/ptodata/2/ina/peTUS_COMB.seq:*
... /cgn2_6/ptodata/2/ina/peTUS_COMB.seq:*
... /cgn2_6/ptodata/2/ina/peTUS_COMB.seq:*
... /cgn2_6/ptodata/2/ina/packfilesI.seq:*
GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                       Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                 383533 seqs, 122816752 residues
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Maximum Match 100%
Listing first 45 summaries
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1 ATGATCGAAACATACAACCA...
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Gapop_10.0 , Gapext 1.0
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Maximum DB seq length: 200000000
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Perfect score:
Sequence:
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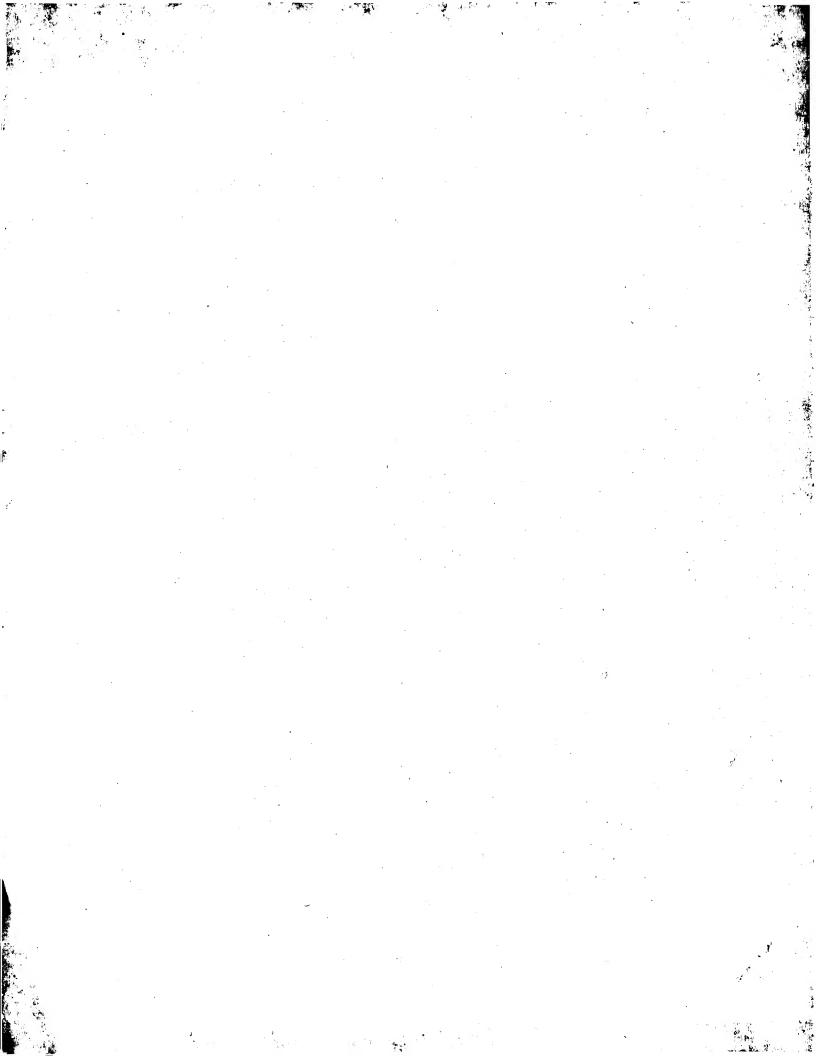
ALIGNMENTS

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APPLICANT: Spriggs, Melanie
APPLICANT: Srinivasan, Subhashini
TITLE OF INVENTION: Methods of Preparing Soluble, Oligomeric
TITLE OF INVENTION: Proteins
                                                                                                                                                                                                                                                                                                                                           Apple Operating System 7.1 ft Word for Apple, Version 5.1a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              USSN 08/107,353
                                                                                                                                                                                                                                                                                                                                                                                                     US/08/446,922
                                                                                                                                                                            3: Immunex Corporation
51 University Street
Sequence 3, Application US/08446922
Patent No. 5716805
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Perkins, Patricia A
REGISTRATION NUMBER: 34,693
REFERENCE/DOCKET NUMBER: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                 Apple Macintosh
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (206)587-0430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Perkins, Patricia
                                                                                                                                                                                                                                                                                                           Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (206)233-0644 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     . 786 base pairs
nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: 08-13-93
                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS: single
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MEDIUM TYPE: Floppy
                                                                                                                                          NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                   Microsoft
                                                                                                                                                                                                                                                                                                                                               OPERATING SYSTEM:
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STRAIN: CI
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Sequence 1, Sequence 15,

Sequence

Sednence



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SEQ ID NO 10646

LENGTH: 57649

CORGANISM: Homo sapiens

US-10-105-299-10646

Guery Match

Best Local Similarity 54.0%; Pred. No. 32;

Best Local Similarity 54.0%; Pred. No. 32;

Matches 75; Conservative 0; Mamarches 64; Indels 0; Gaps. (0; Matches 75; Conservative 0; Mamarches 64; Indels 0; Gaps. (0)

QY 161 AGAGGAACTACATCATGAAGATTATTCATAAAAACTAAAAGAGATCAAACA 220

QY 221 AAGAGAACTAAACCTTCATGAAGATTATTCATAAAAACTAAAAAGAAGACAATTGAAGAC 280

QY 221 AAGAGAACTATATCATCATCAACTAATAAAAACAAGAAGACCAATTGAAGAC 280

QY 221 AAGAGAACTATAACCTT 299

QY 281 TTGTCAAGGATTAAACGTT 299

COURTEAL INFORMATION: Human Secreted Proteins

FILE REPERENCE: PSSSO

COURTENT PAPLICATION: Human Secreted Proteins

FILE REPRENCE: PSSSO

COURTENT PAPLICATION: HUMAN SECRETED PSSSO

COURTENT PAPLICATION: HUMAN SECRETED PSSSO

COURTENT PSSSO

COURTENT
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Ouery Match
Best Local Similarity 54.0%; Pred. No. 32;
Matches 75; Conservative 0; Mismatches 64; Indels 0; Gaps 0;
Oy 161 AAGAGGAAGTAAAACCTTCATGAAGATTTGTATTCATAAAAAGCTAAAGATGCAACA 220
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TYPE: DNA CORGANISM: Homo sapiens US-10-105-299-14274 Search completed: May 30, 2002, 05:45:48 Job time: 17122 sec

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221 AAGGAGAAGGATCTTTATCCTTGCTGAACTGTGAGAGATGAGAAGGCAATTTGAAGACC 280
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                                                                                                                                                              255 GGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTAAACAAAGAAGAAG 314
                                                                                                                                                                                                467 GCAAATGAACTAGACATGCCAACAGCTTGGGGAAGAGAATTCCATACTGAAAGAAGAGCA 408
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                                                                                         Length 637;
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CURRENT FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 15197
Prior Application removed - See File Wrapper or Palm SOFTWARE: Patentin Ver. 2.0
                                                                                                                            97;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               64;
                                                                                         DB 6;
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                                                                                       Score 36.8; DE
Pred. No. 9;
0; Mismatches
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Pred. No. 32;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Rosen, et. al
TITLE OF INVENTION: Human Secreted Proteins
FILE REFEBENCE: PS950
CURRENT APPLICATION NUMBER: US/10/105,299
CURRENT FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 15197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10646, Application US/10105299
GENERAL INFORMATION:
APPLICANT: Rosen, et. al
TILLE OF INVENTION: Human Secreted Proteins
FILE REFERENCE: PS950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 9151, Application US/10105299
GENERAL INFORMATION:
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Best Local Similarity 49.55
Matches 95; Conservative
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US-10-105-299-9151
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               ; ORGANISM: Human US-10-027-632-217676
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PRIOR APPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR APPLICATION NUMBER: US 60/198,676
PRIOR FILING DATE: 2000-04-20
PRIOR APPLICATION NUMBER: US 60/193,483
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR APPLICATION NUMBER: US 60/185,218
PRIOR PILING DATE: 2000-03-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR PILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/166,358
PRIOR APPLICATION NUMBER: US 60/166,358
PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR FILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FASTSED for Windows Version 4.0
SEQ ID NO 217675
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ FD NO 217676
LENGTH: 637
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Best Local Similarity 49.5
Matches 95; Conservative
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US-10-027-632-217675
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TYPE: DNA

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TITLE OF INVENTION: Colon and Colon Cancer Associated Polynucleotides and Polypept FILE REFERENCE: PA005P1
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                                                                                                                                      263 igcitaittaacagaigaaaagaagtaaatittacaaigaagccacagitictaaittai 322
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GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827,129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
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    Score 37.2; DB 6; Length 544; Pred. No. 6.9; 0; Mismatches 73; Indels
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Pred. No. 7.
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PRIOR APPLICATION NUMBER: US-60/157,137
PRIOR PILING DATE: 1999-09-29
PRIOR PILING DATE: 1999-09-29
PRIOR PLING DATE: 1999-09-29
PRIOR PLICATION NUMBER: US-60/163,280
PRIOR FILING DATE: 1999-11-03
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Best Local Similarity
Matches 81; Conserv
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SOFTWARE: Patentin Vel
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US-10-027-632-217675/c
                                    Local Similarity
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US-10-106-698-1404
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                 Query Match
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GENERAL INFORMATION:
APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
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Pred. No. 16;
3; Mismatches 106; Indels
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PRIOR FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR PILIAG DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/156,358
PRIOR APPLICATION NUMBER: US 60/146,002
PRIOR PILING DATE: 1999-09-09
PRIOR FILING DATE: 1999-08-09
NUMBER OF SEQ ID NOS: 325720
SOFTWARE: FASTEED for Windows Version 4.0
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CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
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PRIOR FILING DATE: 2000-07-12
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APPLICATION NUMBER: US 60/193,483
FILING DATE: 2000-03-29
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PPLICATION NUMBER: US 60/185,218
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Best Local Similarity 48.18;
Matches 101; Conservative
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LOCATION: (1)...(12231
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                                                                                                               2158 AGGATAGGTTATATTAGATTTCATATGCTTTATTTACTAAGGCAGAGAAATGGGCAATAC 2099
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                       121 CTTTTTGCTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT 180
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                                                                                            ATGAAAATTTTTATGTATTTACTTACTGTTTTTCTTATCACCCAGATGATTGGGTCAGCA 120
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
FILE REFERENCE: 108827.129
CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
  1 ATGATCGAAACATACAACCATCTCCCCGATCTGCGCCCACTGGACTGCCCATCAGC
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Pred. No. 0.88;
0; Mismatches 149; Indels
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PRIOR FILING DATE: 2000-07-12
PRIOR PELING DATE: 2000-07-12
PRIOR PELING DATE: 2000-07-20
PRIOR FILING DATE: 2000-04-20
PRIOR PELICATION NUMBER: US 60/193, 483
PRIOR APPLICATION NUMBER: US 60/195, 218
PRIOR APPLICATION NUMBER: US 60/165, 218
PRIOR FILING DATE: 2000-02-24
PRIOR FILING DATE: 1999-11-23
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/167, 363
PRIOR FILING DATE: 1999-11-23
PRIOR PRIOR APPLICATION NUMBER: US 60/167, 369
PRIOR FILING DATE: 1999-10-28
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SOFTWARE: FastSEQ for Windows Version 4.0
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Best Local Similarity 46.8
Matches 131; Conservative
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US-10-027-632-111808/c
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ORGANISM: Human
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LENGTH: 2865
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                                                    APPLICANT: Wang, David G.
TITLE OF INVENTION: Identification and Mapping of Single Nucleotide
TITLE OF INVENTION: Polymorphisms in the Human Genome
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TITLE OF INVENTION: Identification and Mapping of Single Nucleotide TITLE OF INVENTION: Polymorphisms in the Human Genome TITLE OF INVENTION: Polymorphisms in the Human Genome CURRENT APPLICATION NUMBER: US/10/027,632 CURRENT FILING DATE: 2002-04-30
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Pred. No. 1.7;
                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/10/027,632
CURRENT FILING DATE: 2002-04-30
PRIOR PAPLICATION NUMBER: US 60/218,006
PRIOR FILING DATE: 2000-07-12
PRIOR PAPLICATION NUMBER: US 60/218,616
PRIOR PAPLICATION NUMBER: US 60/193,483
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 1900-03-24
PRIOR FILING DATE: 1999-11-23
PRIOR PLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-11-23
PRIOR APPLICATION NUMBER: US 60/167,363
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
PRIOR FILING DATE: 1999-09-28
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PRIOR FILING DATE: 2000-07-12
PRIOR PLICATION NUMBER: US 60/198,676
PRIOR PILING DATE: 2000-04-20
PRIOR FILING DATE: 2000-03-29
PRIOR FILING DATE: 2000-03-29
PRIOR APPLICATION NUMBER: US 60/185,218
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Sequence 233265, Application US/10027632
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Best Local Similarity 48.99
Matches 107; Conservative
                                                                                                                                                                 FILE REFERENCE: 108827.129
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                                                                            148 AACTTGGTAACCCTGGAAAATGGGAAACAGCTGACCGTTAAAAGACAAGGACTCTATTAT
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Pred. No. 6.8e-29;
); Mismatches 17;
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CURRENT APPLICATION NUMBER: US/09/875,453A
CURRENT FILING DATE: 2002-05-06
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Sheppard, Liana T.
Lim, Moon Young
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Michelotti, Emil E
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Thomas, Rita L.
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Pred. No. 5.2e-82;
); Mismatches 85;
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SOFTWARE: FastSEQ for Windows Version 4.0
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PRIOR APPLICATION NUMBER: US 60/220,057
PRIOR FILING DATE: 2000-07-21
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APPLICANT: Dunn, II, Robert T
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US-09-911-904-131
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US-09-911-904-131
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Best Local S
Matches 422
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Pred. No. 2.2e-169;
0; Mismatches 27;
                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: LUKAShev, Matvey
TITLE OF INVENTION: Hematology/Immunology Array
FILE REFERENCE: CLON-006CIP15
CURRENT APPLICATION NUMBER: 05/09/442,384B
CURRENT FILING DATE: 1999-11-17
PRIOR APPLICATION NUMBER: 09/053,375
PRIOR FILING DATE: 1998-03-31
NUMBER OF SEQ ID NOS: 830
                                                                                                                                                                                                                                                                                                                                               Sequence 457, Application US/09442384B
GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                92.8%;
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Matches 756; Conservative
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; ORGANISM: homo sapiens
US-09-442-384B-457
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1 ATGATCGAAACATACAACCTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC
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Pred. No. 1e-142; 0; Mismatches 103;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Chenchik, Alex
APPLICANT: Bibilashvilli, Robert
TITLE OF INVENTION: Nucleic Acid Arrays
FILE REPERENCE: CLON-006
CURRENT APPLICATION NUMBER: US/09/053,375B
CURRENT FILING DATE: 1998-08-31
NUMBER OF SEO ID NOS: 1543
SOFTWARE: FastSEO for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 821, Application US/09053375B GENERAL INFORMATION:
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86.8%;
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Best Local Similarity 86.8
Matches 680; Conservative
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(without alignments)
1310.273 Million cell updates/sec
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783
1 ATGATCGAAACATACAACCA......TTGGCTTACTCAAACTCTGA 783
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Perfect score:
Sequence:
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/cgn2_6/ptodata/1/pna/US60_NEW_COMB.seq: pna/US10 NEW

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Pending_Patents_NA_New:

Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysis of the total score distribution.

						SUMMARIES		
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Result No.	ult No.	Score	Query Match	Length	DB	. QI	Description	æ
;	-	726.8	1 2	1816	5	US-09-053-375B-226	Sequence 22	26, Ap
	7	726.8	92.8	1816	2	9-47	Sequence 45	-
	m	618.2		1250	S	US-09-053-375B-821		21, A
	4	371	47.4	508	Ŋ	US-09-911-904-131	Sequence 131	31, AF
	2	155.8	19,9	2395	Ŋ	19-875-	Sequence 9,	
υ	9	41.6		2865	9	US-10-027-632-111808		1808,
O	7	39.8	5.1	637	9	US-10-027-632-233265 .	(7)	233265,
U	8	39.2	5.0	1223197	9	US-10-027-632-179264	a	179264
	6	37.2	4.8	544	9	US-10-027-632-45446	4	45446,
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υ	12	36.8	4.7	637	-	US-10-027-632-217676		17676,
	13	36.6	4.7	57649	-	US-10-105-299-9151		9151, 1
	14	36.6	4.7	57649		US-10-105-299-10646		10646,
	15	36.6	4.7	57649		US-10-105-299-14274		1274,
Ö	16	36.4	4.6	2580		32-2		58659,
	17	36.2	4.6	2430		US-10-105-694-1	Sequence 1,	App.
	18	36.2	4.6	2430		US-10-105-695-1	Sequence 1,	, App.
	19	9	4.6	2430		US-10-106-014-1		, App.
υ	20		4.6	559		-10-		78162,
ņ	21		4.6	559		US-10-027-632-78163	-	m
	22	36	4.6	573		US-10-027-632-263768		263768
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	24	35.6	4.5	. 501		7-632-		9002,
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61 100

40 atgategaaacatacaaaccaaactteteeeegatetgeggeeaetggaetgeeeateage

1 ATGATCGAAACATACAACCAAACTTCTCCCCGATCTGCGGCCACTGGACTGCCCATCAGC

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Gaps

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Length 1816; Indels

Score 726.8; DB 5; Pred. No. 2.2e-169; 0; Mismatches 27;

92.8%;

Query Match 92.8 Best Local Similarity 96.2 Matches 756; Conservative

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В Qy

121

CTITITIGCTGTGTATCTTCATAGAAGATTGGATAAGGTCGAAGAGGAAGTAAACCTTCAT

GAAGATTTTGTATTCATAAAAAGCTAAAGAGATGCAACAAAAGGAGAAGGATCTTTATCC

TTGCTGAACTGTGAGGAGATGAGAAGGCAATTTGAAGACCTTGTCAAGGATATAACGTTA 300

AACAAAGAAGA - - - GAAAAAAGAAAACCACTTTGAAATGCAAAAAGGTGATCAGAATCCT

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